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OM nucleic - nucleic search, using sw model
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Sequence 39928, A	Sequence 2765, Ap	Þ	sequence 2766, Ap	Sequence 41384, A	Sequence 41384, A	Sequence 41384, A	Sequence 41392, A	Sequence 41392, A	Sequence 41392, A	Sequence 819, App	Sequence 818, App	•	`	•		010	010,		Describeron	7)

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RESULT 2
PCT-US01-01333-819
; Sequence 819, Application PC/TUS0101333
; GENERAL INFORMATION:
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; Sequence 818, Application PC/TUS0101333
; GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 819
               APPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008PCT
CURRENT APPLICATION NUMBER: PCT/US01/01333
CURRENT FILING DATE: 2001-01-14
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 1198
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008BCT
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SOFTWARE: PatentIn Ver. 2.0
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CURRENT FILING DATE: 2001-01-14
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Similarity 88.8%;
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PCT-US01-01354-30829
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US-09-764-891-8151
US-09-764-905-26661
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PCT-US01-01339-6894
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US-09-950-083-6997
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US-10-092-399-39928
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Sequence 6894, Ap
Sequence 8151, Ap
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Sequence 10742, A
Sequence 10746, A
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                                         Query Match
Best Local Similarity
Matches 87; Conserv
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LENGTH: 98
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SEQ ID NO 818
LENGTH: 98
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
CURRENT APPLICATION NUMBER: US/09/764,860
CURRENT FILING DATE: 2001-01-17
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Best Local Similarity 88.8%;
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                                                                                                                                                                                                                    Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1198
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Best Local
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
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                                                                                                                                                         TYPE: DNA
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4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 CCACCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGG 98
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Similarity 88.8%;
                                                                                                                                                                                                                                                                                                                                                                     Application US/09764860
                                           Conservative
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                                                              88.8%;
                                       0; Mismatches 11;
                                                            Score 80.4; DB 30;
Pred. No. 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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RESULT 5
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GENERAL INFORMATION
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TITLE OF INVENTION: Nucleic Acids, Proteins,
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RAPPLICATION NUMBER: 60/226,868
RAPPLICATION SAME CONTROL OF CON
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                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/228,924 FILING DATE: 2000-08-30 APPLICATION NUMBER: 60/224,518
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APPLICATION NUMBER: 60/234,223
FILING DATE: 2000-09-21
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                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/224,519 FILING DATE: 2000-08-14
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FILING DATE: 2000-11-17
APPLICATION NUMBER: 60/236,327
FILING DATE: 2000-09-29
APPLICATION NUMBER: 60/241,785
                                                                                   APPLICATION NUMBER: 60/241,809 FILING DATE: 2000-10-20 APPLICATION NUMBER: 60/249,299 FILING DATE: 2000-11-17
                                                                                                                                                                                 APPLICATION NUMBER: 60/220,964 FILING DATE: 2000-07-26
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/246,474
PRIOR FILING DATE: 2000-11-08
PRIOR APPLICATION NUMBER: 60/246,532
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DR FILLING DATE: 2000-08-14
DR APPLICATION NUMBER: 60/227,182
DR FILLING DATE: 2000-08-22
DR APPLICATION NUMBER: 60/225,214
DR APPLICATION NUMBER: 60/225,214
DR FILLING DATE: 2000-08-14
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APPLICATION NUMBER: 60/
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APPLICATION NUMBER: 60/244,617
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                                                                     FILING DATE: 2000-09-27
APPLICATION NUMBER: 60/230,438
FILING DATE: 2000-09-06
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                                                                                                                                                      APPLICATION NUMBER: 60/235,836
                                                  APPLICATION NUMBER: 60/215,135
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60/237,037

60/239,937 60/239,935

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                                                 4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGTGGTCTCGAACTCTTTGACCTCGGGTGAT 4149
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OR APPLICATION NUMBER: 60/249,264
OR FILING DATE: 2000-11-17
OR APPLICATION NUMBER: 60/249,214
OR FILING DATE: 2000-11-17
OR APPLICATION NUMBER: 60/249,297
OR FILING DATE: 2000-11-17
OR APPLICATION NUMBER: 60/232,400
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                             1 GTAGAGACGGGGTTTCACCATGTTGGCCAGGCTGATCTCGAACTCCTGACCTCAGGTGAT 60
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APPLICATION NUMBER:
FILING DATE: 2000-0
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FILING DATE: 2000-11-17
APPLICATION NUMBER: 60/249,217
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APPLICATION NUMBER: 60/249,207
FILING DATE: 2000-11-17
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APPLICATION NUMBER: 60/249,212
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                                                                                                                  87;
                                                                                                                             Similarity
                                                                                                            Conservative
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Pred. No. 2;
                                                                                                                       Pred. No.
                                                                                                       Mismatches
                                                                                                                                        DB 39;
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US-10-074-095-819; Sequence 819, Applic; GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-09-29
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PRIOR TITING DATE: 2000-08-14
PRIOR APPLICATION NUMBER: 60/226,868
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/216,647
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PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: 60/214,886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 09/764,860 PRIOR FILING DATE: 2001-01-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/179,065 PRIOR FILING DATE: 2000-01-31
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FILING DATE: 2000-10-20
APPLICATION NUMBER: 60/244,617
FILING DATE: 2000-11-01
                                                     APPLICATION NUMBER: 60/236,327
FILING DATE: 2000-09-29
APPLICATION NUMBER: 60/241,785
                                                                                                                     APPLICATION NUMBER: 60/241,809
FILING DATE: 2000-10-20
APPLICATION NUMBER: 60/249,299
FILING DATE: 2000-11-17
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FILING DATE: 2000-09-27
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FILING DATE: 2000-08-14
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FILING DATE: 2000-08-14
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                                                                                                                                                                                                                                                      60/224,519
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2000-11-17

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PRIOR PRIOR
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DR FILING DATE: 2000-08-14

DR APPLICATION NUMBER: 60/235,836

DR FILING DATE: 2000-09-27

DR APPLICATION NUMBER: 60/230,438

DR FILING DATE: 2000-09-06

DR APPLICATION NUMBER: 60/215,135

DR FILING DATE: 2000-06-30

DR APPLICATION NUMBER: 60/225,266

DR FILING DATE: 2000-08-14

DR APPLICATION NUMBER: 60/249,218
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REAPPLICATION NUMBER: 60/237,040
REFILING DATE: 2000-10-02
OR APPLICATION NUMBER: 60/240,960
REFILING DATE: 2000-10-20
REPPLICATION NUMBER: 60/239,935
REPPLICATION NUMBER: 60/239,935
REPPLICATION NUMBER: 60/239,935
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APPLICATION NUMBER: 60/246,532
APPLICATION NUMBER: 60/249,216
APPLICATION NUMBER: 60/249,216
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APPLICATION NUMBER: 60/237,038
FILING DATE: 2000-10-02
APPLICATION NUMBER: 60/236,370
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FILING DATE: 2000-10-13
APPLICATION NUMBER: 60/241,787
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APPLICATION NUMBER: 60/225,214
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APPLICATION NUMBER: 60/227,182
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APPLICATION NUMBER: 60/249,210
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APPLICATION NUMBER: 60/246,474
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APPLICATION NUMBER: 60/237,037
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APPLICATION NUMBER: 60/231,413
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APPLICANT: Rosen et
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               TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PC004PCT CURRENT APPLICATION NUMBER: PCT/US01/01354 CURRENT FILING DATE: 2001-03-17
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
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                                                                                           APPLICANT: Human Genome Sciences, Inc. et al
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TITLE OF INVENTION:
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PRIOR APPLICATION NUMBER: 60/220,963
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                               FILING DATE: 2000-08-30
APPLICATION NUMBER: 60/224,518
FILING DATE: 2000-08-14
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FILING DATE: 2000-07-07
APPLICATION NUMBER: 60/225,267
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APPLICATION NUMBER: 60/235,834
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APPLICATION NUMBER: 60/226,868
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OR APPLICATION NUMBER: 60/224,519
OR FILING DATE: 2000-08-14
OR APPLICATION NUMBER: 60/220,964
OR FILING DATE: 2000-07-26
OR APPLICATION NUMBER: 60/241,809
OR FILING DATE: 2000-10-20
OR FILING DATE: 2000-10-20

APPLICATION NUMBER: 60/249,299 FILING DATE: 2000-11-17

APPLICATION NUMBER: 60/236,327

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APPLICATION NUMBER: 60/229,509
FILING DATE: 2000-09-05

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OR APPLICATION NUMBER: 60/237,038
OR FILING DATE: 2000-10-02
OR APPLICATION NUMBER: 60/236,370
OR APPLICATION NUMBER: 60/236,802
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OR APPLICATION NUMBER: 60/237,037
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OR FILING DATE: 2000-10-20
OR APPLICATION NUMBER: 60/239,935
OR FILING DATE: 2000-10-13
OR APPLICATION NUMBER: 60/239,937
OR FILING DATE: 2000-10-13
OR APPLICATION NUMBER: 60/241,787

OR APPLICATION NUMBER: 60/249,216
OR FILING DATE: 2000-11-17
OR APPLICATION NUMBER: 60/249,210
OR FILING DATE: 2000-11-17
OR APPLICATION NUMBER: 60/226,681
OR APPLICATION NUMBER: 60/225,759
OR APPLICATION NUMBER: 60/225,759
OR FILING DATE: 2000-08-14

FILING DATE: 2000-11-08
APPLICATION NUMBER: 60/246,532
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FILING DATE: 2000-09-05
APPLICATION NUMBER: 60/236,367
FILING DATE: 2000-09-29
APPLICATION NUMBER: 60/237,039
FILING DATE: 2000-10-02

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OR FILING DATE: 2000-09-01
OR APPLICATION NUMBER: 60/229,287
OR FILING DATE: 2000-09-01
OR APPLICATION NUMBER: 60/229,513
OR FILING DATE: 2000-09-05
OR APPLICATION NUMBER: 60/231,413

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OR APPLICATION NUMBER: 60/229,344
OR FILING DATE: 2000-12-08
OR APPLICATION NUMBER: 60/229,344
OR FILING DATE: 2000-09-01
OR PILING DATE: 2000-09-07
OR FILING DATE: 2000-09-25
OR APPLICATION NUMBER: 60/229,343
OR APPLICATION NUMBER: 60/229,343
OR APPLICATION NUMBER: 60/229,343

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APPLICATION NUMBER: 60/225,268 FILING DATE: 2000-08-14 APPLICATION NUMBER: 60/236,368

APPLICATION NUMBER: 60/251,856 FILING DATE: 2000-12-08

FILING DATE: 2000-09-29

PRIOR PRIOR PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/244,617
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FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/241,785

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PCT-US01-01354-41384
Sequence 41384, Application PC/TUS0101354;
GENERAL INFORMATION:
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SEQ ID NO 41384
LENGTH: 99
TYPE: DNA
                                                                    Matches
                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US01/01354
CURRENT FILING DATE: 2001-03-17
NUMBER OF SEQ ID NOS: 42506
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CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 42506
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TYPE: DNA
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               PRIOR APPLICATION NUMBER: 60/233,065 PRIOR FILING DATE: 2000-09-14 PRIOR APPLICATION NUMBER: 60/232,398
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nl Similarity 88.8%;
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Pred. No. 2;
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RESULT 13
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                                                             PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 2000-09-29
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PRIOR APPLICATION NUMBER: 60/214,886
PRIOR FILING DATE: 2000-06-28
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR FILING DATE: 2000-07-11
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PRIOR FILING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: 60/180,628
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 APPLICATION NUMBER: 60/244,617 FILING DATE: 2000-11-01
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FILING DATE: 2000-09-29
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FILING DATE: 2000-08-14
APPLICATION NUMBER: 60/225,213
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APPLICATION NUMBER: 60/225,266
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APPLICATION NUMBER: 60/249,218
FILING DATE: 2000-11-17
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Pred. No. 2.5;
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RESULT 14

US-10-092-399-41384

Sequence 4184, Application US/10092399

Sequence 4181, Application US/10092399

GENERAL INFORMATION:

APPLICANT: Rosen et al.

TITLE OF INVENTION: NUCLEIC Acids, Proteins, and Antibodies

FILE REFERENCE: PC004C1

CURRENT APPLICATION NUMBER: US/10/092,399

CURRENT FILING DATE: 2002-03-07

NUMBER OF SEQ ID NOS: 42506

Prior Application removed - See File Wrapper or Palm

SOFTMARE: Patentin Ver. 2.0

SEQ ID NO 41384
Search completed: January 4, 2003, 20:26:35 Job time: 8810 secs
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PCT-US01-01324-2765
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GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-01-14
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC002PCT
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                                                                          4150 CCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACAAG 4187
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TYPE: DNA
ORGANISM: Homo sapiens
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Perfect score:
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Copyright (c) 1993 - 2003 Compug
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   /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*
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/cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
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US-09-513-999C-19853
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US-09-513-999C-20263
US-09-513-999C-17082
US-09-513-999C-18226
US-09-513-999C-2027453
US-09-513-999C-16174
US-09-513-999C-16266
US-09-513-999C-17641
US-09-513-999C-17641
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	ESULT 2 S-09-513-999C-17594 S-09-513-999C-17594 Sequence 17594, Application US/09513999C Sequence 17594, Application US/09513999C GENERAL INFORMATION: APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y. APPLICANT: Giordano, J.Y. APPLICANT: CONTRON: Expressed Sequence Tags and Encoded Human FILE REFERENCE: 59.US2.REG CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24	Duery Match 1.1%; Score 75.4; DB 5; Length 96; Best Local Similarity 87.2%; Pred. No. 0.081; Matches 82; Conservative 0; Mismatches 12; Indels 0; Gaps 0 4110 TGTTGGCCGGGCTGGTCTGAACTCTTGACCTCGGTGATCCACCCAC	LENGTH: 90 TYPE: DNA ORGANISM: Homo sapienS FEARTURE: NAME/KEY: misc_feature LOCATION: 86 OTHER INFORMATION: n=a, g, c or t S-09-513-999C-16529	APPLICANT: Girdano, J.Y. APPLICANT: Girdano, J.Y. APPLICANT: Girdano, J.Y. APPLICANT: Girdano, J.Y. FILE OF INVENTION: Expressed Sequence Tags and Encoded Human FILE REFERENCE: 59 JUS2. REG CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26 NUMBER OF SEQ ID NOS: 36681 SOUTWARE: Patent.pm SEQ ID NO 16529	ESULT 1 S-09-513-999C-16529 Sequence 16529, Application US/095 GENERAL INFORMATION: GENERAL INFORMATION: DEPLICANT: Dumas Milne Edwards, J	36 60 0.9 95 5 US-09-513-999C-18970 Sequence 18970 37 59.6 0.9 95 5 US-09-513-999C-25357 Sequence 25357 38 59.2 0.9 88 5 US-09-513-999C-23355 Sequence 25357 40 59 0.9 91 5 US-09-513-999C-21342 Sequence 21342 40 58.8 0.9 99 5 US-09-513-999C-18125 Sequence 213236 41 58.8 0.9 92 5 US-09-513-999C-18125 Sequence 213236 42 58.4 0.9 92 5 US-09-513-999C-23356 Sequence 22336 43 58.2 0.9 91 5 US-09-513-999C-23375 Sequence 23375 44 57.4 0.9 90 5 US-09-513-999C-21158 Sequence 23158	27 61.4 0.9 89 5 US-09-513-999C-17626 Sequence 17626, 28 61.4 0.9 97 5 US-09-513-999C-35981 Sequence 35981, 29 61 0.9 77 5 US-09-513-999C-35981 Sequence 36945, 29 61 0.9 93 6 US-10-170-385-494 Sequence 494, 4 30 61 0.9 96 5 US-09-513-999C-17020 Sequence 17020, 31 61 0.9 96 5 US-09-513-999C-18968 Sequence 18968, 32 61 0.9 96 5 US-09-513-999C-18968 Sequence 18968, 32 61 0.9 96 5 US-09-513-999C-18968 Sequence 18968, 33 61 0.9 96 5 US-09-513-999C-18968 Sequence 18968, 34 60.4 0.9 82 5 US-09-513-999C-18840 Sequence 18840, 35 60.4 0.9 92 5 US-09-513-999C-18840 Sequence 18840, 35 60.4 0.9

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US-09-513-999C-15261
Sequence 15261, Application US/09513999C
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-513-999C-19855
SOFTWARE: Patent.pm
SEQ ID NO 15261
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                               APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59 USZ. REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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SOFTWARE: Patent.pm
SEQ ID NO 19855
LENGTH: 98
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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SOFTWARE: Patent.pm
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                                                                                                                                                                                                                                                                                                                                            61 TTCCTGAGTAGCTGGGATTACAGGCGCACACCACCAT 97
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Similarity 84.5%;
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Pred. No. 0.16;
0; Mismatches 15; Indels
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US-09-513-999C-15268
                                                                                        Sequence 15268, Application US/09513999C
GENERAL INFORMATION:
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                                                                 APPLICANT: Dumas Milne Edwards, J.B.
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Best Local Similarity
FILE REFERENCE: 59.US2.REG
   APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
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APPLICANT:
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LENGTH: 97
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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OTHER INFORMATION: w=a or t
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Local Similarity 83.78;
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                                                      Duclert, A.
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                US-09-513-999C-19833/c; Sequence 19833, Application US/09513999C; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
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GENERAL INFORMATION:
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Best Local (
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SEQ ID NO 19652
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SOFTWARE: Patent.pm
SEQ ID NO 15268
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Best Local :
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins
FILE REFERENCE: 59.US2.REG
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Local Similarity 81.6%;
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Duclert,
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Pred. No. 0.25;
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SSOFTWARE: Patent.pm
SEQ ID NO 19825
LENGTH: 95
TYPE: DNA
ORGANISM: Homo sapiens
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Sequence 20263, Application US/09513999C
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
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LENGTH: 93
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Best Local
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PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
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RESULT 12
US-09-513-999C-20264
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                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and
FILE REFERENCE: 59.US2.REG
SOFTWARE: Pate
SEQ ID NO 20264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 19012
LENGTH: 89
TYPE: DNA
                                                                                                                                                                                                                                                     Sequence 20264, Application US/09513999C
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LENGTH: 97
                                                    CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                      NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487
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APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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Local Similarity 84.4%;
es 76; Conservativo
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Local Similarity 85.2%;
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                                                                                                                                      Query Match 1.0%;
Best Local Similarity 82.2%;
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SEQ ID NO 17082
LENGTH: 93
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and
FILE REFERENCE: 59.US2.REG
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                                                                                                                                                                                                                NAME/KEY: misc_feature LOCATION: 34
                                                                                                                                                                                                                                                  OTHER INFORMATION: s=g or FEATURE:
                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature LOCATION: 17
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61
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GGACTACAGGTGCCCACGACCATGCCCGGC 90
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Pred. No. 1;
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RESULT 14
US-09-513-999C-34460/c
; Sequence 34460, Application US/09513999C
; GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.

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RESULT 15
US-09-513-999C-18226/c
; Sequence 18226, Application US/09513999C
; GENERAL INFORMATION:
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; LOCATION: 7:
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US-09-513-999C-18226
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Search completed: January Job time : 330 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 1.0%; Score 65.6; DB 5; Length 89; Best Local Similarity 84.1%; Pred. No. 1.5; Matches 74; Conservative 0; Mismatches 14; Indels
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SEQ ID NO 34460
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Best Local
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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
FEATURE:
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                                                                                                        5234 CCTGCCTCAGCCTCCCAAAGTGCTGGGA 5261
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Local Similarity 83.0%;
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1.1 99 13 1.1 95 9 1.1 82 14 1.1 99 13 1.1 94 10	Score
BI017753 PM3-ET02 AA669571 ac18dl0.s T81021 yd25b03.s1 BG977835 MR2-cr011 AW339643 he15e10.:	Description
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T82910 T82910 B0446872 AA828120 F24101 A2757066 AA494010	AA486/52 AW072612 BG674517 BF130198	A116816/ AF219090 BG108795	AA077606 AA729064 AA923035	BG029559 AA665078	AA454805 AA180452	AA649287 AZ576255	BF542018 F28887	AA053038 BH770807 AA280198	AA608742 AA457423	F30221 F26219	AA486800 AF149649	861 474
7d39d0; 7d39d0; 7d39d0; 7d71a; 0d71a; 1SPD10; 6 ew08; ng60b;	52 ab1/b03 612 xa37d09 517 6026209 198 6018182	AF219 HRPE	nw22f09. ok82a01.	6022967 nu76b10.		ns12h07.s	10 0	ZI/IeI/	ae56g04. aa86b02.	SPD20571 SPD13614	ab19a06.r AF149649	507 RC2-CT0 905 xy21d11

ALIGNMENTS

	MEDLINE	JOURNAL	TITLE				AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	RESULT 1 BI017753
Laboratory of Cancer Genetics Ludwig Institute for Cancer Research Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP, Brazil Tel: +55-11-2704922	20202663 Contact: Simpson A.J.G.	sequence tags Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)	Simpson, A.J. Shotgun sequencing of the human transcriptome with ORF expressed	,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and	Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare	Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,	Dias Neto, E., Garcia Correa,K., Verjovski-Aimerda,S., birones,M.K.,	1 (bases 1 to 99)	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human.	EST.	BI017753.1 GI:14421824	BI017753	77-160401-003-h09_1 ET0277	BIO17753 99 bp mRNA linear EST 13-JUN-2001	

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VERSION
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                                                                                                                                                            Seq
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1 (bases 1 to 95)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
                                                                                                                                                                                                   Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL ; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA669571 95 bp mrNA linear EST 20-NOV-19: ac18d10.s1 Stratagene ovary (#937217) Homo sapiens cDNA clone IMAGE:856819 3' similar to contains Alu repetitive element;, mrNA
                                                                                                                                                              IMAGE Consortium (info@image.llnl.gov) fo
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                         Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                    Washington University School of Medicine
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM3&t2=PM3-ET0277-
160401-003-h09_l&t3=2001-04-16&t4=1)
Seq primer: puc 18 forward
High quality sequence stop: 99.
                                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1997)
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/clone_lib="Stratagene ovary (#937217)"
/sex="female"
                                              /clone="IMAGE:856819"
                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tissue mRNA and cDNA amplification were performed under low stringency conditions."
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                                                                                                                                      Location/Qualifiers
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/clone_lib="ET0277"
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                     Email: est@watson.wustl.edu
Insert Size: 725
High quality sequence stops: 73 Source: IMAGE Consortium,
clone is available royalty-free through LLNL; contact the
Consortium (info@image.llnl.gov) for further information.
Insert Length: 725 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R. The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Other_ESTs: yd25b03.rl
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                                                                                                                                                                                                                                                                                                                                            Seq primer: -21m13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1995)
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               /dev_stage="49 year old"
/lab_host="SOLR cells (Kanamycin resistant)"
/note="Organ: ovary; Vector: Bluescript SK; Site_1: EcoRI;
/note="Organ: ovary; Vector: Bluescript SK; Site_1: Cligo dr.
Total ovary tissue, normal, caucasian. Average insert
size: 0.8 kb; Uni-ZAP XR Vector; -5' adaptor sequence: 5'
GAATTCGGCACGAG 3' -3' adaptor sequence: 5'
CTCGAGGTTTTTTTTTTTTTTTTTTTTTTTTTTTT 3'"
double-stranded cDNA was ligated to Eco RI adaptors
                                                                                                                                                 /sex="male"
                                                                                                                                                                       /clone_lib="Soares fetal liver spleen lNFLS"
                                                                                                                                                                                             /db_xref="taxon:9606"
/clone="IMAGE:109229"
                                                                                                                                                                                                                                       /db_xref="GDB:464846"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 CAAAGTTTTGGGATTACAGG 81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l (bases 1 to 99)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence.
BG977835
                                                                                                                                                                                                                                                                                                                                                                                                        Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
Project. This entry can be seen in the following URL
                                                                                                                                                                                                                                                                                                                                        High
                                                                                                                                                                                                                                                                                                                                                           (http://www.ludwig.org.br/scripts/gethtml2.pl?ti=MR2&t2=MR2-CI0128-110101-016-f07_1&t3=2001-01-11&t4=1)
Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shotgun sequencing of the human transcriptome with ORF expressed
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Mammalia;
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MR2-CI0128-110101-016-f07_1 CI0128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence tags
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Location/Qualifiers
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                                                                              /note="Organ: colon_ins; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of
                  tissue mRNA and cDNA amplification were performed under low stringency conditions." 34\ c\qquad 28\ g\qquad 21\ t\qquad 3\ others
                                                                                                                                                                                                    /db_xref="taxon:9606"
/clone_lib="CI0128"
/dev_stage="Adult"
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                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
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Pred. No. 0.061;
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RESULT 5
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Best Local Similarity
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61
                                                                                                                                                                                     Local Similarity
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                       TCTCCTGCCTCAGCCTCCCAAAGTGCTGGGATTA 5264
                                                                                                                       GTAGAGCCAGGGTCTCACTATGTTGCCCAGGCAGGTCTTGAACTCCTGGCCTCAAGTGAT 5230
CCTCCCACCCGGCTTCCCAAAATGCTGGGATTA 94
                                                                               GTAGAGATGTGGTCTCACTATGTTGCCCCAGGCTGGTCTCGAACTCCTGGGCTCAAGTGAT 60
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Genome Sequencing Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ggapbs-r@mail.nih.gov
Tissue Procurement: Elisabeth Paietta, Jonathan D. Licht, M.D.,
Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life
Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Possible reversed clone: polyT not found Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 94)
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                                                                                                                                                                                                                                                                  /note="Organ: whole blood; Vector: pCMV-SPORT6; Site_1: Sall; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies."

28 c 24 g 24 t
                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:2919114"
/clone_lib="NCI_CGAP_CML1"
/tissue_type="myeloid cells, 18 p
rearrangement positive, includes
myeloid blast crisis"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                   1.1%;
85.1%;
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81.8%;
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Pred. No. 0.
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Pred. No. 0.058;
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RESULT 6
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                                                                           AUTHORS
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l (bases 1 to 99)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P..., Tongapagai C.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chen, H., Huo, Y., Patel, S., Zhu, X., Swift-Scanlan, T., Reeves, R.H., DePaulo, R. Jr., Ross, C.A. and McInnis, M.G.
Gene identification using exon amplification on human chromosome 18q21: implications for bipolar disorder 18q21: Psychiatry 5 (5), 502-509 (2000)
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AF149606 Human chromosome 18q21
genomic clone 3h1, DNA sequence.
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Johns Hopkins University School of Medicine
600 N. Wolfe Street, Baltimore, MD 21287, US
Email: hc@welchlink.welch.jhu.edu
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26 c
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                  GI:7957200
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25 g 25 t 1 others
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                                                                                                                                            JOURNAL
                                                                                                                                                                                                          AUTHORS
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Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Ludwig Institute for Prudente 109, 4 andar,
Emmert-Buck, M.D., Ph.D.
Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: Life Technologies,
cDNA Library Arrayed by: Greg Lennon, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                     xy21d11.x1 NCI_CGAP_Ut4 Homo sapiens cDNA clone IMAGE:2853813 3' similar to contains Alu repetitive element;contains element MER9 repetitive element; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=RC2-CT0320-281
199-012-b11_1&t3=1999-11-28&t4=1)
Seq_primer: puc 18 forward
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                                                                                                                                       Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                           Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
                                                                                             Email: cgapbs-r@mail.nih.gov
                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                             Tumor Gene Index
                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stringency conditions."
17 c 33 g
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/db_xref="taxon:9606"
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82.5%;
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                                                                                                                                                                                    Project (CGAP),
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                        Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                   Seq primer: Promega -21m13
                                                                                                            Email: est@watson.wustl.edu
Insert Size: 174
                                                                                                                                                             Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                             Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                            On Oct 18, 1995 this sequence version replaced gi:1026289
                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yu68f10.s1 Weizmann Olfactory Epithelium Homo sapiens cDNA clone IMAGE:238987 3' similar to contains Alu repetitive element; contains MER22 repetitive element; mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/TLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 98)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Average insert size 1.48 kb. Life Technologies catalog #: 11542-016"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sall; Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="serous papillary carcinoma, high grade,
pooled tumors"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="IMAGE:2853813"
/clone_lib="NCI_CGAP_Ut4"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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81.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 69.2; DB 10; Pred. No. 0.17;
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 CCCAGGCCGGAGTGCAATGGCGCGACCTCAGCTCACTGCAACCCCTGCCTCCCAGGTTCA 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                     High quality sequence stop: 97.
                                                                                                                                   Seq primer: -28m13 rev1 ET from Amersham
                                                                                                                                                  The vector to vector length is 165
Insert Length: 1338 Std Error: 0
                                                                                                                                                                                                                                                                  Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                             Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin, J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ab19a06.rl Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:841234 5' similar to contains Alu repetitive element;contains element MER22 repetitive element;, mRNA sequence.
                                                                                                                                                                                           Putative full length read
                                                                                                                                                                                                  This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                               Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                     Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                     White, Y., Wylie, T., Watersto
WashU-NCI human EST Project
                                                                                                                                                                                                                                           Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                              4444 Forest Park Parkway, Box 8501, St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 99)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
/sex="male"
           /clone_lib="Stratagene lung (#937210)"
                                   /clone="IMAGE:841234"
                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jocation/Qualifiers
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/organism="Homo sapiens"
/db_xref="GDB:3864371"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="Weizmann Olfactory Epithelium"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="IMAGE:238987"
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                                                                                                                                                                                                                                                                                            Louis, MO 63108
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                                                                       3956 CTGGCGTGCAGTGGGATCTGGGTTCACTGCAACCTCCGCCTCTTGGGTTCAAGCGAT 4015
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1 CTGGAGTGCAGTGGCGGATCTCGGCTCACTGCAGCCTTGACCTCCTGGGCTCAAGCAAT 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chen.H., Huo,Y., Patel,S., Zhu,X., Swift-Scanlan,T., Reeves,R.H., DePaulo,R. Jr., Ross,C.A. and McInnis,M.G. Gene identification using exon amplification on human chromosome 18q21: implications for bipolar disorder 18q21: psychiatry 5 (5), 502-509 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF149649.1 GI:8485987
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Chen H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Johns Hopkins University School of Medicine
600 N. Wolfe Street, Baltimore, MD 21287, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Psychiatry and Behavioral Sciences
                                                                                                                                                                                              Similarity
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                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="72 years"
//ab_host="SOLR cells (kanamycin resistant)"
//ab_host="SOLR cells (kanamycin resistant)"
//note="Organ: lung: Vector: pBluescript SKr; Site_1: EcoRI / Note="Organ: lung: Vector | Disposed | Dispose
                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Human chromosome 18q21 from exon-trapping"
/note="Bacterial artificial chromosome library prepared in
/note="Bacterial artificial chromosome library prepared in
pBACe3.6 vector via DNA partially digested with a
combination of EcoRI and EcoRI Methylase for library
segments 1&2 or either MboI or DpnII for library segments
segments 1&2 or either MboI or DpnII for library segments
3&4. Size selected DNA was cloned into the pBACe3.6 vector
at the EcoRI sites for library segments 1&2 (plates 1-480)
or the BamHI sites for library segments 3&4 (plates
                                                                                                                                                                                                                                                                                                                                      481-1056). http://www.chori.org/bacpac/"
31 c 28 g 22 t
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/db_xref="taxon:9606"
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                                                                                                                                                               0;
                                                                                                                                                                                              Score 66.6; DB Pred. No. 0.45;
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/note="Vector: pcDNAII (Invitrogen); Site_1: BStXI;
/note="Vector: pcDNAII (Invitrogen); Site_1: BStXI;
Site_2: NotI; The library was constructed by G.
Lanfranchi. This library is not subtracted nor normalized.
The first strand cDNA was primed with a biotinylated
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/db_xref="taxon:9606"
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                                                                 cDNA clone s4000051B06, mRNA sequence.
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RESULT 14
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            Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins M., Hultman, M., Kucaba, T., Lacy, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry, Meg, J., Trevaskis, E.,
                                                                                                                                                                                                                                                                                                          T62174 87 bp mRNA linear EST 14-FEB-1 yb96c02.r1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:79010 5' similar to contains Alu repetitive element;, mRNA
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CRIBI Biotechnology Centre
                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M.
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(bases 1 to 87)
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/clone="s4000051B06"
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Best Local :
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5253 GTGCTGGGATTACAGGTGTG 5272
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                                                                                                                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 90)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone is available royalty-free through LLNL; contact th consortium (info@image.llnl.gov) for further information. Insert Length: 447 Std Error: 0.00 seg primer: m13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996) 97044478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ae56g04.s1 Stratagene lung carcinoma 937218 Homo se IMAGE:950934 3' similar to contains Alu repetitive
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High qality sequence stops: 55 Source: IMAGE Consortium,
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                                                                        Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St.
                                                                                                                              Unpublished (1997)
Contact: Wilson RK
                                                                                                                                                                                   White,Y., Wylie,T., WaterstowashU-NCI human EST Project
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est@watson.wustl.edu
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/note-"Organ: lung; Vector: pBluescript SK: Site_1: EcoRI
/site_2: XhoI; Cloned unidirectionally. Primer: Oligo
dT. normal lung. Average insert size: 1.0 kb; Uni-ZAP XR
Vector: -5' adaptor sequence: 5' GATTCGGCAGGAG 3' -3'
adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTTTTTTTTTTTT
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/db_xref="GDB:482627"
/db_xref="taxon:9606"
/clone="TMAGE:79010"
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Search completed: January 4, 2003, 17:56:55 Job time : 5563 secs
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ORIGIN
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Best Local Similarity 83.1%;
Matches 74; Conservative
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                                                 62 AGGTGATCTGCCTGCCTCGGCCTCCCAAA 90
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                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                               Score 65; DB 9; Length 90; Pred. No. 0.85; 0; Mismatches 15; Indels
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Listing first 45 summaries
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    pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 99
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1: /cgn2_6/ptodata/1,

2: /dgn2_6/ptodata/1,

3: /cgn2_6/ptodata/1,

4: /cgn2_6/ptodata/1,

5: /cgn2_6/ptodata/1,

6: /cgn2_6/ptodata/1,
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6672
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14935.382 Million cell updates/sec
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
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/cgn2_6/ptodata/1/ina/backfiles1.seq:*
    Length
    DB
                                                US-09-357-740-15
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US-08-3450-673c-91
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US-08-988-128-2
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15. Appl
17. Appl
17. Appl
18. Company
18. Company
19. Appl
19.
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RESULT 2 US-09-357-740-14 ; Sequence 14, Application US/09357740 ; Patent No. 6348596 ; GENERAL INFORMATION: ; APPLICANT: Lee, Linda G. ; APPLICANT: Graham, Ronald J. ; APPLICANT: Mullah, Khairuzzaman B. APPLICANT: Haxo, Francis T. ; TITLE OF INVENTION: ASYMMETRIC CYANINE DYE QUENCHERS ; FILE REFERENCE: 9584-007	Query Match Query Matches Query Ma	RESULT 1 US-09-357-740-15 US-09-357-740-15 Sequence 15, Application US/09357740 Patent No. 6348596 Patent No. 6348596 GENERAL INFORMATION: APPLICANT: Graham, Ronald J. APPLICANT: Graham, Ronald J. APPLICANT: HANO, Francis T. TITLE OF INVENTION: ASYMMETRIC CYANINE DYE QUENCHERS FILE REFERENCE: 9584-007 CURRENT APPLICATION NUMBER: US/09/357,740 CURRENT FILING DATE: 1999-07-20 EARLIER APPLICATION NUMBER: US/09/357,740 CURRENT FILING DATE: 1998-01-23 NUMBER OF SEQ ID NOS: 22 SOFTWARE: PATENTIN VET: 2.0 SEQ ID NO 15 LENGTH: 75 TYPE: DNA ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Oligonucleotide US-09-357-740-15	28 42.4 0.6 60 2 US-08-340-426D-59 Sequence 59, Appl 29 42.4 0.6 60 3 US-08-450-673C-59 Sequence 59, Appl 20 42.4 0.6 60 3 US-08-545-860D-71 Sequence 71, Appl 32 42.4 0.6 60 5 PCT-US94-04496-71 Sequence 71, Appl 32 42.4 0.6 60 5 PCT-US95-17111A-59 Sequence 59, Appl 34 40.8 0.6 85 2 US-08-340-426D-92 Sequence 92, Appl 35 40.8 0.6 85 2 US-08-340-426D-92 Sequence 92, Appl 37 40.8 0.6 85 2 US-08-340-426D-92 Sequence 92, Appl 38.6 0.6 85 5 PCT-US95-17111A-79 Sequence 92, Appl 40 38.6 0.6 78 2 US-08-340-426D-70 Sequence 92, Appl 40 38.6 0.6 78 2 US-08-32-766A-3 Sequence 37, Appl 41 38.6 0.6 78 2 US-08-340-426D-70 Sequence 70, Appl 41 38.6 0.6 78 2 US-08-340-673C-70 Sequence 70, Appl 42 38.2 0.6 43 2 US-08-988-128-7 Sequence 70, Appl 50 50 50 50 50 50 50 50 50 50 50 50 50
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            Sequence 91, Application US/08454557C Patent No. 5830670 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/461,697
CURRENT FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 466
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 130
LENGTH: 75
                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Barney, Shawn
APPLICANT: Thomas, Mary Beth
APPLICANT: Thomas, Mary Beth
APPLICANT: Portbury, Stuart D.
APPLICANT: Puranam, Kasturi
APPLICANT: EVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
TITLE OF INVENTION: AND TREATING CONDITIONS, DISORDERS, OR DISEASES INVOLVING
TITLE OF INVENTION: CELL DEATH
FILE REFERENCE: 10001-005-999
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APPLICANT: COGENT NEUROSCIENCE, Inc
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GENERAT TOTAL
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LENGTH: 75
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EARLIER FILING DATE: 1998-01-23
NUMBER OF SEQ ID NOS: 22
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                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
 APPLICANT:
                                                                                                                                                                     4177
                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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                                                                                                                                   14 GGGATTACAGGCAT 1
                                                                                                                                                                                            Local
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                                                                                                                                                      GGGATTACAAGCGT 4190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69;
                                                                                                                                                                                                                                                                                  Similarity
de la Monte, Suzanne
                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                  0.9%;
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                                                                                                                                                                                                                                                                              Score 59.6; DB 4; Length 75; Pred. No. 5.9e-05;
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Pred. No. 2.7e-06;
                                                                                                                                                                                                                                                                Mismatches
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US-08-340-426D-91
; Sequence 91, Application US/08340426D
; Patent No. 5948634
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Best Local Similarity
                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: de la Monte, S
APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/340,426D
                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                              TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection TITLE OF INVENTION: of Alzheimer's Disease NUMBER OF SEQUENCES: 121
                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4167 CCAAAGTGCTGGGATTACAAGCGTG 4191
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TELECOMMUNICATION INFORMATION:
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NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,
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APPLICATION NUMBER: US
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CITY: Washington
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FILING DATE:
                                                                                                                                     ZIP:
                                                                                                                                                   COUNTRY:
                                                                                                                                                                       STATE:
                                                                                                                                                                                  CITY: Washington
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                                                                                                                                     20005-3934.
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20005-3934
                                                                                                                                                                   D.C.
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EDNESS: both
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SYSTEM: PC-DOS/MS-DOS
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14-NOV-1994
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87.1%;
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Pred. No. 0.00031;
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; Sequence 91, Application US/08450673C
; Patent No. 5948888
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Query Match 0.8%; Score 56.6; DB 2; Best Local Similarity 87.1%; Pred. No. 0.00031; Matches 74; Conservative 0; Mismatches 9
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                                                                                                                                                               TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
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REGISTION NUMBER: 36,203
REFERENCE, DOCKET NUMBER: 066
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: Ludwig, Steven R.
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                         NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
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Local Similarity 87.1%;
nes 74; Conservative
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                                                                                                             TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                     FILING DATE: 30 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/OFILING DATE: 30-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
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EDNESS: both
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                                                                                                                                                                                                                           (202) 371-2600
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 Mismatches

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                                                                                                                                                                                                                                                                    0609.3840004
                                          DB 2; Length 84;
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                                                                                                                                                                  Query Match
Best Local S
                                                                                                                                                Matches
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INFORMATION FOR SEQ ID NO: $
SEQUENCE CHARACTERISTICS:
LENGTH: 84 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Ludwig, Steven R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/3
FILING DATE: 14-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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                                                                                        4167 CCAAAGTGCTGGGATTACAAGCGTG 4191
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                                                                                                                                                                                                                                     TYPE: nucleic acid STRANDEDNESS: both TOPOLOGY: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US95/17111A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                Watch 0.8%; Score 56.6; DB 5; Length 84; Local Similarity 87.1%; Pred. No. 0.00031;
                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Ludwig, Steven REGISTRATION NUMBER:
59 CCAAAGTGCTGGGATTACAAGCGTG 83
                                                                     1 CCATGTTCATCAGGCTGGTGTCGAACTCCTGACCTC--GTGATCCGCCCGCCTCAGCCTC 58
                                                                                                                                              74; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RY: U.S.A.
20005-3934
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                                                                                                                                                                                                                                                            both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neural Thread Protein Gene Expression and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08/340,426
                                                                                                                                                                                                                                                                                                                                                                                                                             36,203
                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                             0609.3840002
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                                                                                                                                                      Indels
                                                                                                                                                          2
                                                                                                                                                        Gaps
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; Sequence 1, Application US/08988128
; Patent No. 5994505

GENERAL INFORMATION:

APPLICANT: Ting, Jenny Pan-Yung APPLICANT: Chin, Keh-Chin

RESULT 8 US-08-988-128-1

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US-08-988-128-2
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                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 88 base pairs
                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                 APPLICANT: Ting, Jenny Pan-Yung
APPLICANT: Chin, Keh-Chin
TITLE OF INVENTION: No. 5994505el Forms of Class II MHC
TITLE OF INVENTION: Transactivator (CIITA)
                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (919) 854-1400
TELEFAX: (919) 854-1401
                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                STREET:
CITY: F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          187 CAAG 190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: No. 5994505el Forms of Class II MHC TITLE OF INVENTION: Transactivator (CIITA)

CORRESPONDENCE ADDRESS:
APPLICATION NUMBER: US/C
FILING DATE: 10-DEC-1997
                                                                                                                                                 COUNTRY:
                                                                                                                                                                    STATE:
                                                                                                                                                                                          ADDRESSEE: Myers Bigel Sibley & Sajovec, P.A. STREET: P.O. Box 37428
                                                                                                                                                                                                                                                                                                                                                                                                                                         85 CAAG 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Biswas, Sorojini J. REGISTRATION NUMBER: 39,111 REFERENCE/DOCKET NUMBER: 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/O FILING DATE: 10-DEC-1997 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 0.8%;
Local_Similarity 92.2%;
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                                                                                                                                   27627
                                                                                                                                          Raleigh
No. 5994505th Carolina
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No. 5994505th Carolina
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                                                                                              Floppy disk
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Pred. No. 0.00043;
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; GENERAL INFORMATION:
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                                                                                                                                                     RESULT 11
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APPLICANT: Swensen, Jeff
TITLE OF INVENTION: A 14 KILOBASE DELETION IN THE PROMOTER REGION OF BRCA1;
TITLE OF INVENTION: IN A BREAST CANCER FAMILY
FILE REFERENCE: 2318-197
CURRENT APPLICATION NUMBER: US/09/056,762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-988-128-2
                                                                                             Sequence 151, Application US/09461697 Patent No. 6277974
                                                                                                                                                                                                                                                                                                                                                                 Best
APPLICANT: COGENT NEUROSCIENCE, Inc. APPLICANT: Lo, Donald C. APPLICANT: Barney, Shawn APPLICANT: Thomas, Mary Beth
                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 60/043,116
EARLIER FILING DATE: 1997-04-09
NUMBER OF SEQ ID NOS: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                    4010 AGCGATTCTTCTGC 4023
                                                                                                                                                                                                                                                                               3950 CCCAGGCTGCAGTGCAGTGCGATCTGGGTTCACTGCAACCTCCGCCTCTTGGGTTCA 4009
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LENGTH: 88 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                              75 CTCAGGCTGGAGTGCAGTGGTGGTGGTTGGCTCACTGCAACCTCTGCCTCTCGGATTGA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      127 GATTCCTACACAATGCGTTGCCTGCCTCCACGCCCTGCTGGGTCCTACCTGTCAGAGCCC 186
                                                                                                                                                                                                    15 AGTGATTCTTGTGC 2
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                                                                                                                                                                                                                                                                                                                                                          Match 0.8%; Local Similarity 83.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           85 CAAG 88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Biswas, Sorojini J. REGISTRATION NUMBER: 39,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                            62;
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) 854-1401
- NO: 2:
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92.2%;
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                                                                                                                                                                                                                                                                                                                                   Score 54.8; DB 3; Length 75; Pred. No. 0.00076; 0; Mismatches 12; Indels
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Pred. No. 0.00043;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-461-697-151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-454-557C-60
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CURRENT FILING DATE: 1999-12-14
NUMBER OF SEQ ID'NOS: 466
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 151
LENGTH: 75
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%;
Best Local Similarity 83.8%;
Matches 62; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 60, Application US/08454557C Patent No. 5830670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: de la Monte, S
APPLICANT: Wands, Jack R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Katz, Lawrence C.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING TITLE OF INVENTION: AND TREATING CONDITIONS, DISORDERS, OR ITLE OF INVENTION: CELL DEATH FILE REFERENCE: 10001-005-999
                                                                                                                                                             TELEFAX: (202) 371-254
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
TELECHEONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Portbury, Stuart D. APPLICANT: Puranam, Kasturi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4042 TGGGACTACAGGCA 4055
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                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection TITLE OF INVENTION: of Alzheimer's Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 30-MA' CLASSIFICATION: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1100 New CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                            LENGTH: 60 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Sterne, Kessler, Goldstein & Fox P.L.L.C 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 U.S.A.
                                                                                    both
                                                                                                                                                                                                                                                                                                                                                                         30-MAY-1995
0.8%;
91.7%;
                                                                                                                                                                                                         2540
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Pred. No. 0.00076;
0; Mismatches 12;
                                                                                                                                                                                                                                                                     0609.3840003
Score 52; DB 2;
Pred. No. 0.0031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12;
                   Length 60;
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US-08-340-426D-60
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                                                                                                                                                                                    Patent No. 5948888 GENERAL INFORMATION:
                                                                                                                                                                                                Sequence 60, Apprint No. 5948888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                        APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                           4121 CTGGTCTCGAACTCTTGACCTCGGGTGATCCACCCACCTCAGCCTCCCAAAGTGCTGGGA 4180
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LENGTH: 60 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection TITLE OF INVENTION: of Alzheimer's Disease NUMBER OF SEQUENCES: 121
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            CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
COUNTRY: U.S.A.
                                                ADDRESSEE: Sterne, Kessler, G
STREET: 1100 New York Avenue,
                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/0 FILING DATE: 14-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 20005-3934
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C STREET: 1100 New York Avenue, Suite 600
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EDNESS: both
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91.7%;
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                                                               Kessler, Goldstein &
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                                                Suite 600
                                                                Fox P.L.L.C
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                                                             ATTORREY/AGENT INFORMATION:

NAME: Ludwig, Steven R.

REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609.3840002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2540
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 60:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCT-US95-17111A-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%;
Best Local Similarity 91.7%;
Matches 55; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 60, Application PC/TUS9517111A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ. ID NO: 60:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/450,673C
FILING DATE: 30-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
21P: 20005-9934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
SEQUENCE CHARACTERISTICS:
LENGTH: 60 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/340,426
FILING DATE: 14-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene Expression and
TITLE OF INVENTION: Detection of Alzheimer's Disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Ludwig, Steven R. REGISTRATION NUMBER: 36,203
                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US95/17111A FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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Job time : 144 secs

Search completed: January 4, 2003, 17:59:29

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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Match
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      US-09-761-288-47

US-09-764-860-882

US-09-864-761-25274

US-09-736-969A-5

US-09-74-877-3656

US-09-764-904-89

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US-09-764-847-1866
US-09-764-847-1867
  US-09-764-860-608
US-09-764-860-609
US-09-764-869-1767
US-09-764-869-1992
US-09-764-869-1993
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US-09-764-860-819
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Sequence 1865, Ap
Sequence 1867, Appl
Sequence 47, Appl
Sequence 682, Appl
Sequence 5274, A
Sequence 5, Appli
Sequence 3656, Ap
Sequence 79, Appl
Sequence 80, Appl
Sequence 81, Appl
Sequence 607, App
Sequence 608, App
Sequence 609, App
Sequence 1767, Ap
Sequence 1992, Ap
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Sequence 819, App
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-764-878-2	-764-877-2	69-1	-869-1	64-877-3	64-869-2	US-09-764-869-2102	61-288-3	64-877-	47-	64-847-115	-860-9	64	-764-887-	61-288-4	1-288-4	64-855-	-764-877-	-764-860-	64-887	64-860	64-860-105	-764-860-	4-860-	US-09-764-869-1281	-69
Sequence 204, App	`	-	-	e 3775,	Sequence 2105, Ap	Sequence 2102, Ap	2	`		•	Sequence 962, App	Sequence 2198, Ap	Sequence 559, App	`	, 1	25	ω (8)	836,	54,	Sequence 1053, Ap	'n	Sequence 643, App	42,	28	Sequence 1719, Ap

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RESULT 1
US-09-764-860-818
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
CURRENT APPLICATION NUMBER: US/09/764,860
CURRENT FILING DATE: 2001-01-17
                                                                                                 Sequence 819, Application US/09764860 Patent No. US20020094953A1 GENERAL INFORMATION:
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                                      SENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prior application data removed - consult PALM or NUMBER OF SEQ ID NOS: 1198
                   CURRENT APPLICATION NUMBER: US/09/764,860
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TYPE: DNA
ORGANISM: Homo sapiens
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    2001-01-17
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Pred. No. 9.1e-06;
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; Prior application data removed - co
; NUMBER OF SEQ ID NOS: 1198
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 819
; LENGTH: 98
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-860-819
; ORGANISM: Homo sapiens US-09-764-847-1866
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US-09-764-868-1444/c
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                       CURRENT APPLICATION NUMBER: US/09/764,847
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 2003
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1866
LENGTH: 98
TYPE: DNA
TYPE: DNA
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APPLICANT: ROSen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PT332
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
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                                                                                                                                                                                                       APPLICANT: Rosen et al. TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PC009
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TYPE: DNA
ORGANISM: Homo sapiens
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Local Similarity 87.4%;
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Pred. No. 9.1e-06;
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; ORGANISM: Homo sapiens
US-09-764-847-1867
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LENGTH: 98
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Best Local
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                                                                                                                                                             TITLE OF INVENTION: No. US20020065405A1el Polypeptides and Nucleic Acids Encoding FILE REFERENCE: 15966-638
CURRENT APPLICATION NUMBER: US/09/761,288
CURRENT FILING DATE: 2001-01-16
PRIOR APPLICATION UNDER: 60/177,839
PRIOR FILING DATE: 2000-01-25
PRIOR APPLICATION NUMBER: 60/176,134
PRIOR APPLICATION NUMBER: 60/176,134
PRIOR FILING DATE: 2000-01-14
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APPLICANT: Prayaga, Sudhirdas
APPLICANT: Taupier, Raymond J
APPLICANT: Mishra, Vishnu
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                                                                                                 PRIOR APPLICATION NUMBER: 60/175,989
PRIOR FILING DATE: 2000-01-13
PRIOR APPLICATION NUMBER: 60/218,324
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APPLICANT:
                                                                                    PRIOR FILING DATE:
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC009
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FILING DATE:
                      APPLICATION NUMBER: 60/178,191
                                        APPLICATION NUMBER: 60/220,253
FILING DATE: 2000-07-24
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Pred. No. 6.2e-05;
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Pred. No. 6.2e-05;
0; Mismatches 14;
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; TYPE: DNA
'; ORGANISM: Homo sapiens
US-09-761-288-47
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LENGTH: 96
TYPE: DNA
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Best Local Similarity 85.4%;
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Best Local
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PRIOR FILING DATE: 2000-01-26
PRIOR APPLICATION NUMBER: 60/220,590
PRIOR FILING DATE: 2000-07-25
NUMBER OF SEQ ID NOS: 95
                                                                                                                                                                                                                                  Sequence 25274, Application US/09864761 Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Rosen et al. TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PC008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/764,860 CURRENT FILING DATE: 2001-01-17
                  CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
                                                                                          TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                 APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K
APPLICANT: Chen, Wensheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0
                                                                          FILE REFERENCE: Aeomica-X-1
PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
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                                                                                                                                   Hanzel, David K.
Chen, Wensheng
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  2000-02-04
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Pred. No.
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
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PRIOR APPLICATION NUMBER: US 60/234,687
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                                                                      5235 CTGCCTCAGCCTCCCAAAGTGCTGGGATTACAG 5267
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APPLICATION NUMBER:
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APPLICATION NUMBER: PCT/US01/00668
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APPLICATION NUMBER: US 09/632,366
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CTGCCTTGGCTTCCCAAAATGCTACAATTACAG
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NE EXPRESSED IN LUNG, SIGNAL = 3.8

NE EXPRESSED IN PLACENTA, SIGNAL = 3

NE EXPRESSED IN HELA, SIGNAL = 4.2

NE EXPRESSED IN HELA, SIGNAL = 4.2

NE EXPRESSED IN FETAL LIVER, SIGNAL = 5.2

NE EXPRESSED IN FETAL LIVER, SIGNAL = 5.2
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N: EXPRESSED IN HEART, SIGNAL = 4.5

N: EXPRESSED IN LUNG, SIGNAL = 3.8

N: EXPRESSED IN PLACENTA, SIGNAL = 3.8

N: EXPRESSED IN BRAIN, SIGNAL = 4.2

N: EXPRESSED IN HELA, SIGNAL = 4.2

N: EXPRESSED IN ADULT LIVER, SIGNAL = 4.2

N: EXPRESSED IN FETAL LIVER, SIGNAL = 4.2

N: EXPRESSED IN FETAL LIVER, SIGNAL = 4.2
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                                                                                                                                                                                                                                                                                                                      Score 70.6; DB 10; Pred. No. 0.00045;
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RESULT 9 US-09-736-969A-5

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Sequence 5, Application US/09736969A Patent No. US20020068302A1 GENERAL INFORMATION:

Lu, Peter

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                                 GENERAL INFORMATION:
APPLICANT: ROSen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC005
CURRENT APPLICATION NUMBER: US/09/764,877
CURRENT FILING DATE: 2001-01-17
CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                 Sequence 3656, Application US/09764877 Patent No. US20020147140A1
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Best Local
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Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 4031
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SOFTWARE: PatentIn Ver. 2
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PRIOR APPLICATION NUMBER: US 60/196,460
PRIOR FILING DATE: 2000-04-11
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PRIOR FILING DATE: 2000-04-11
PRIOR APPLICATION NUMBER: US 60/196,267
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PRIOR APPLICATION NUMBER: US 60/240,503
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PRIOR FILING DATE: 2000-01-14
PRIOR APPLICATION NUMBER: US 60/182,296
PRIOR FILING DATE: 2000-02-14
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                                                                                                                                                                                                                                                                                                                                                                                          4017 CTTCTGCTTCAGCCTCCCGAGTAGCTGGGACTACAGG 4053
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                                                                                                                                                                                                                                                                                                                                          62 CTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTGTAGG 98
                                                                                                                                                                                                                                                                                                                                                                                                                                   2 TGGAGTGCAGTGACGCGATCTCGGCTCGCTACAACCTCTGCCTCCCGGGTTCAAGTGATT 61
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82.5%;
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Pred. No. 0.00063;
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; LENGTH: 90
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-904-80
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US-09-764-904-80/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-877-3656
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SEQ ID NO 79
LENGTH: 90
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LENGTH: 90
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 Best Local Similarity
                     Query Match
                                                                                                                                                                                                                                                                                                                           Patent No. US20020173454A1
                                                                                                                                                                                                                                                                                                                                            Sequence 80,
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA122
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CURRENT FILING DATE: 2001-01-17
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Best Local Similarity 87.4%;
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Local Similarity 85.4%;
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Pred. No. 0.0012;
Score 68.2; DB 9
Pred. No. 0.0012;
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                 DB 9;
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                 Length 90;
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                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/764,860
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1198
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 607
LENGTH: 90
TYPE: DNA
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC008
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LENGTH: 90
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity 85.4%;
Matches 76; Conservative
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA122
CURRENT APPLICATION NUMBER: US/09/764,904
CURRENT FILING DATE: 2001-01-17
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29 AAGTGATCTGCCTGCCTCGGCCTCCCAAA 1
                                                                        89 ATTTTTAGTAGAGACAGGGTTTCACCATGTTGGCCAGGCTGGTCTCGAACTCCTGACCTC 30
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-860-608
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US-09-764-860-608/c
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Search completed: January
Job time : 192 secs
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LENGTH: 90
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Best Local Similarity
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                                                                                                           4143 GGGTGATCCACCCACCTCAGCCTCCCAAA 4171
                                                                      29 AAGTGATCTGCCTGCCTCCCAAA 1
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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6672
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

TITLE	REFERENCE AUTHORS	SOURCE	ACCESSION VERSION	RESULT 1 AX197482/ LOCUS DEFINITION		C 44 C 45				c 37 38	36	ω ω π 4-		c c 31			25	c 24 25		c 21		18	c 16		c 13		c 10	ဝ ထ	7 0	าเก	.e. u		7	Result No. S	
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 Hobbs, H.H.,
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Analysis of the LDL-receptor gene of a patient with familial hypercholesterolemia (FH) revealed the deletion of exon 5 resulting hypercholesterolemia (FH) hotween repetitive Alu sequences of
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Human low density lipoprotein receptor gene (LDLR),
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Pred. No. 0
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                                    G00-119-362"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTGCTGGGATTACAG 75
                                                                     Hobbs,H.H., Brown,M.S., Goldstein,J.L. and Russell,D.W. Deletion of exon encoding cysteine-rich repeat of low density lipoprotein receptor alters its binding specificity in a subject
Analysis of the LDL-receptor gene of a patient with familial
                        3020025
                                   87008518
                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                   Pluman familial hypercholesterolemia 626-a gene with a deletion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 75)
Lee,L.G., Graham,R.J., Mullah,K.B. and Haxo,F.T.
Non-fluorescent asymmetric cyanine dye compounds useful for quenching reporter dyes
Patent: US 6346596-A 15 19-FEB-2002;
                                                                                                                                       Mammalia; Eutheria; Primates;
                                                                                                                                                                            low density lipoprotein receptor-1; very low density lipoprotein.
                                                                                                                                                                                                                                             exon 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 from patent US 6348596. AR194771
                                                                                                                                                                                                             M14179.1 GI:187101
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                                                    FOTR
                                                                                                                          (bases 1 to 97)
                                         familial hypercholesterolemia
ol. Chem. 261 (28), 13114-13120 (1986)
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                                                                                                                                                   Metazoa; Chordata;
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82.8%;
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Pred. No. 0.00027;
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Pred. No. 0.00022;
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                                                                                                                               Craniata; Vertebrata; Euteleostomi;
Catarrhini; Hominidae; Homo.
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                                      18~a \overline{\phantom{0}3} s \phantom{0}3 c \phantom{0}19~g 20 t \phantom{0}19~p upstream of NheI site; chromosome 19p13.2-p13.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Clean copy of sequence kindly provided by M.Lehrman (22-APR-1987). Individual FH 295 carries two mutant LDL receptor alleles. The allele below was inherited from the father and includes a duplication of exons 2 through 8 that was presumably created by unequal chromosomal crossover involving Alu repeats in introns 1 and 8. The exact site of recombination cannot be determined
                                                                                                                                                                                                                                                                                                                                                                                                                                because the sequences of the Alu repeats in introns 1 and 8 of normal alleles are identical over a span of 26 nucleotides at the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lehrman, M.A., Goldstein, J.L., Russell, D.W. and Brown, M.S.

Duplication of seven exons in LDL receptor gene caused by Alu-Alu
recombination in a subject with familial hypercholesterolemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 90)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human low density lipoprotein receptor mutant gene recombination
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22 a 34 c 23
Chromosome 19p13.2-p13.1.
                                                                                                                                                                                                                                                                                                                                                                                                                recombination site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cell 48 (5), 827-835 (1987)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       recombination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alu repeat; LDL receptor; cell surface protein; crossover;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              M15365.1
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                                                                                    68); putative;
                                                                                                    /note="LDLR duplicated intron 1 (no splice consensus
                                                                                                                                                 /organism="Homo sapiens"
                                                                                                                                                                                                                   /gene="LDLR"
                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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42. .72
                                                                                                                                                                                            /note="LDLR intron
                                                                                                                                                                                                                                                            /gene="LDLR"
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                                                                                                                                                                            . 68
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1.0%; Score 65.6; DB 9;
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Pred. No. 0.00046;
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                                                                               does not fit consensus*
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Sequence 14 from patent US 6348596.
AR194770
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                                                                                                                                                                Lehrman, M.A., Goldstein, J.L., Russell, D.W. and Brown, M.S. Duplication of seven exons in LDL receptor gene caused by Alu-Alu recombination in a subject with familial hypercholesterolemia Cell 48 (5), 827-835 (1987)
                                                                                                                                                                                                                                    Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                      Alu repeat; low density lipoprotein receptor-1; repeat region
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Lehrman, M.A., Gold
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                                                                                                        copy of sequence kindly provided
  Location/Qualifiers
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25 c 18 g
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                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/map="19p13.3"
/note="hgml; G00-119-362"
              /gene="LDLR"
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Best Local Similarity 82.8%;
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                                   3974 ATCTGGGTTCACTGCAACCTCCGCCTC 4000
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Chromosome 19p13.2-p13.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetic and physical mapping of simple sequence repeat containing sequence tagged sites from the human genome Unpublished (1994)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammaîla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 91)

Gerken, S.C., Matsunami, N., Plaetke, R., Albertsen, H., Ballard, L.,
Melis, R., Lawrence, E., Moore, M., Holik, P.R., Carlson, M., Zhao, X.,
Robertson, M., Bradley, P., Elsner, T., Tingey, A., Lalouel, J.-M. and
                                                                                                                                                                                                                                                                                                                                                                                        Cycles Denaturation Annealing C 10 sec. 56 C 10 sec. 72 C 20 sec. 52 C 10 sec. 72 C 20 sec. Mg++: 1.50 r Gel: Acrylamide 7%, Formamide 32%, Urea 34%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Salt Lake City, UT 84112
e-mail: sts@corona.med.utah.edu
Primer A: AGAGGTTGCACTGCACCAA
Primer B: TTTTTCCCCCTCTACTCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             marker; sequence tagged site; tetranucleotide repeat.
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PCR profile:
Initial Denaturation: 94C 300sec
Initial Denaturation Annealing
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Typics 10 sec. 72 C 27
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2160 Eccles Institute of Human Genetics
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/db_xref="taxon:9606"
10...28
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Pred. No. 0.0013;
0; Mismatches 1

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Pred. No. 0.0022;
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                                                                      4111 GTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGATCCACCCCACCTCAGCCTCCCAA 4170
                                                                                                                                                                                                                              source
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                                                                                                                              Local
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             AGTGCTGGGATTACAAGCGTGAGCCACTGCACCGGG 4206
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AGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCCGG
                                                       GTTAGCCAGGATGGTCTCGATCTCCTGACCTC - - ATGATCCACCCGCCTCGGCCTCCCAA 58
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Sequence 48
AX197483
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Sequence 41 from Patent W00151632.
                                                                                                                                                                                                                                                                                  Padigaru,M., Prayaga,S.K., Taupier,R.J., Mishra,V., Tchernev,V.T. Spytek,K.A. and Li,L.
Odorant receptor polypeptides and nucleic acids encoding same
                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 94)
                                                                                                                                                                                                                                                           Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Padigaru,M., Prayaga,S.K., Taupier,R.J., Mishra,V., Tchernev,V.T., Spytek,K.A. and Li,L.
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1 (bases 1 to 94)
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                                                                                                                                                                                                                                                                     receptor polypeptides and nucleic WO 0151632-A 48 19-JUL-2001;
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/db_xref="taxon:9606"
34 c 26 g 1
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/db_xref="taxon:9606"
26 c 34 g 1
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                                                                                                                            Score 62.8; DB 6; Pred. No. 0.0024;
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Pred. No. 0.0024;
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TITLE
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87064495
                                                                                                                                                                                                                                                                                                                                                      Human alpha-galactosidase breakpoint region. M36135
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 80)
                                                                                                                                                                                                                                                                                         Homo sapiens
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Competitive and cooperative functioning of the anterior promoter elements of an Alu family repeat
                                                                                                                                        2160973
                                                                                                                                                        90264427
                                                                                                                                                                                   Alu-rich
                                                                                                                                                                                                Identification of short direct repeats at breakpoints in an
                                                                                                                                                                                                              Alpha-galactosidase A gene rearrangements causing Fabry disease
                                                                                                                                                                                                                              Kornreich, R., Bishop, D.F. and Desnick, R.J
                                                                                                                                                                                                                                                                                                             Human DNA.
                                                                                                                                                                                                                                                                                                                        Fabry disease; breakpoint junction; glycosphingolipid catabolism.
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1 (bases 1 to 94)
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8. >0'
                                                                                                                                                                   Chem. 265 (16), 9319-9326 (1990)
                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
29 c 22 g 1
                                                                                                                                                                                      gene
                                                                                                                      Location/Qualifiers
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23 c 30 g 19 t
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0.9%;
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Score 62.4; DB Pred. No. 0.003;
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ACCESSION
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Best Local Similarity 88.2%;
Matches 67; Conservative
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                                                                         4031 TCCCGAGTAGCTGGGA 4046
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                                                          TCCTGAGTAGCTGGGA 3
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1 (bases 1 to 79)

Schichman,S.A., Caligiuri,M.A., Strout,M.P., Carter,S.L., Gu,Y., Canaani,E., Bloomfield,C.D. and Croce,C.M.

ALL-1 tandem duplication in acute myeloid leukemia with a normal karyotype involves homologous recombination between Alu elements Cancer Res. 54 (16), 4277-4280 (1994)

94320053
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873207
                                                                                                                                                                                                                                                                                                                                                       GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 155288] from the original journal article. This sequence comes from Fig. 3a.

Map location: 11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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/gene="ALL-1"
a 18 c
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/db_xref="taxon:9606"
1. .79
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                                                                                                                                                                      Score 61.6; DB 9; Length 79; Pred. No. 0.0045; 0; Mismatches 9; Indels
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Minimum DB
Maximum DB
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2: /SIDS2/gcgdata/g
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18168.480 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIE

Result	Score	Query Match Length DB	Length		ID	Description
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N	80.4	1.2	98	22	AAS28385	Genomic sequence #
ω	80.4	1.2	99	22	AAK86580	Human immune/haema
4	79.6	1.2	99	22	AAK86572	Human immune/haema
ი 5	78.8	1.2	86	22	AAS32628	Human genomic DNA
თ	78.8	1.2	86	22	AAK89189	Human digestive sy
7	78.8	1.2	98	22	AAK89190	Human digestive sy
ი 8	77.4	1.2	95	22	AAK85116	Human immune/haema
с 9	77.2	1.2	98	22	ABA18411	Human nervous syst

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22222	222	22322	22222	22222222	2222222
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Human Immune/naema Human nervous syst Human nervous syst Human genomic DNA Human immune/haema	immune/ reprodu secrete immune/ immune/	P. P.	Human nervous syst Human genomic DNA Human immune/haema Human immune/haema Human immune/haema	immune/haem testicular nervous sys nervous sys immune/haem immune/haem immune/haem	Human nervous syst Human pancreatic c Human genomic DNA Human reproductive Human reproductive Human immune/haema Human immune/haema Human immune/haema

ALIGNMENTS

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AASZ8384
ID AASZ8384;
XX Human; respiratory entry
ARIONAL ALLERGIC; antigen; respiratory disorder; throat disorder; XX Human; respiratory antigen; respiratory disorder; throat disorder; XX ARIONAL ALLERGIC; anti asthmatic; anti inflammatory; olfactory; XX Respiratory active; ds.
XX ARIONAL ARION
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20-OCT 2000

20-OCT 2000

20-OCT 2000

20-OCT 2000

01-NOV 2000

08-NOV 2000

17-NOV 2000

17-NO
The present invention relates to the isolation of novel human respiratory antigens (AAU17685-AAU17975), and cDNA and genomic sequences encoding for these polypeptides. The sequences of the invention are useful for preventing, treating and/or prognosing disorders related to the respiratory system including throat disorders (e.g. vocal cord paralysis, tonsillitis, and laryngitis),
                                                                                                                                                                                                                                                                                                                     Isolated polypeptide for treating, preventing and/ or prognosing disorders related to the respiratory system including respiratory cancers and also for testing and detection e.g. diagnosis -  \frac{1}{2} \int_{-\infty}^{\infty} \frac{1}{2} \int
                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN GENOME
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28-JUN-2000
30-JUN-2000
707-JUL-2000
11-JUL-2000
11-JU

2000US -0214886 2000US -0216887 2000US -0216880 2000US -0217487 2000US -0217487 2000US -022966 2000US -0224519 2000US -022526 2000US -022526 2000US -022526 2000US -022526 2000US -022527 2000US -022528 2000US -022528 2000US -022528 2000US -022528 2000US -022528 2000US -022528 2000US -022344 2000US -0231244 2000US -0231414 2000US -0231414 2000US -0231414 2000US -0231414 2000US -0231498 2000US -0233497 2000US -023498 2000US -023498 2000US -0235836 2000US -0233680 2000US -0237039 2000US -0237039 2000US -0237039 2000US -023937 2000US

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RESULT 2
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14 - AUG - 2000
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24 - FEB - 2000;
02 - MAR - 2000;
16 - MAR - 2000;
17 - MAR - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; respiratory antigen; respiratory disorder; throat disorder; lung disorder; nose disorder; lung cancer; gene therapy; cytostatic; anti allergic; anti asthmatic; anti inflammatory; olfactory; respiratory active; ds.
                                                                                                                                                                                                                                                                                                                                                                                                        17-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genomic sequence #225 encoding for novel human respiratory antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         lung disorders e.g. pneumonia, allergic disorders e.g. asthma, pleurisy, cystic fibrosis, emphysema, nose disorders and cancers of the respiratory tissues e.g. lung cancer. The polynucleotide sequences of the invention are useful in gene therapy and antisense therapy. human respiratory antigens.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS28385;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GTAGAGACGGGTTTCACCATGTTGGCCAGGCTGATCTCGAACTCCTGACCTCAGGTGAT 60
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                                                                                                                                                                                                                      The present invention relates to the isolation of novel human CC respiratory antigens (AAU17685-AAU17975), and cDNA and genomic CC sequences encoding for these polypeptides. The sequences of the CC invention are useful for preventing, treating and/or prognosing CC disorders related to the respiratory system including throat the CC disorders (e.g. vocal cord paralysis, toosillitis, and laryngitis), CC lung disorders e.g. pneumonia, allergic disorders e.g. asthma, CC lung disorders e.g. pneumonia, allergic disorders e.g. asthma, CC pleurisy, cystic fibrosis, emphysema, nose disorders and cancers of CC the respiratory tissues e.g. lung cancer. The polynucleotide sequences of the invention are useful in gene therapy and antisense therapy. CC AAS28161-AAS28764 represent genomic sequences encoding for novel human respiratory antigens.

CC Note: The sequence data for this patent did not form part of the printed content of the content of the printed content of the content of the content of the printed content in the content of the co
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17 - NOV - 2000;
   Matches
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
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17-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancers and also for testing and detection e.g. diagnosis
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                                                                                                                                  Sequence 98
                               Local Similarity
                                                                                                                                                                                                     ftp.wipo.int/pub/published_pct_sequences.
87;
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2000US-0246532.
2000US-0246609.
2000US-0246610.
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2000US - 024 9297
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2000US-0254097.
2001US-0259678.
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2000US-0249207
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   Conservative
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                                                                                                                                         20 A; 29 C;
                                      1.2%;
88.8%;
                                                                                                                                            27 G; 22 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MS
          0;
                                          Score 80.4; DB 22
Pred. No. 1.5e-06;
              Mismatches
                                                                                 DB 22;
                                                                             Length 98;
                     Indels
                     0;
                     Gaps
                            0;
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RESULT 3
AAK86580
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07-JUN-2000;
28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
14-AUG-2000;

14-AUG-2000;

14-AUG-2000;

14-AUG-2000;

14-AUG-2000;

18-AUG-2000;

22-AUG-2000;

22-AUG-2000;

22-AUG-2000;

23-AUG-2000;

23-AUG-2000;

30-AUG-2000;

01-SEP-2000;

01-SEP-2000;
                                                                                                                                                                                                                                              14-JUL-2000;
26-JUL-2000;
26-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; gene therapy; vaccine; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41392.
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14-AUG-2000;
14-AUG-2000;
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14-AUG-
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11-JUL-2000;
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14-AUG-2000;
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2000US-0224519
2000US-0225213
2000US-0225214
2000US-0225266
2000US-0225267
2000US-0225267
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2000US-0225270
2000US-0225447
2000US-0225757
2000US-0225758
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2000US-0214886

2000US-0215135

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2000US-0217487

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2000US-0217496

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2000US-0205515.
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21-SEP-2000;
21-SEP-2000;
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26-SEP-2000;
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29-SEP-2000;
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08-SEP-2000;
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2000US-0246474.
2000US-0246475.
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2000US-0237037
2000US-0237038
2000US-0237039
2000US-0237040
2000US-0239935
2000US-0239937
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2000US-0246532
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2000US-0241221.
2000US-0241785.
2000US-0241786.
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2000US-0234274
2000US-0234997
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2000US-0235484
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2000US-0246523.
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2000US-0232398
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2000US-0233064.
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2000US-0236370
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2000US-0231242
2000US-0231243
2000US-0231243
2000US-0231413
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2000US-0231414
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RARE REPRESENTATION OF THE PROPERTY OF THE PRO
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17-NOV-2000;
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17-NOV-2000;
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17-NOV-2000;
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2000US-0256719
2000US-0251479
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2000US-0251869
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2000US-0249300.
2000US-0250160.
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2000US-0251030.
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2000US-0249245
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Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis

Disclosure; SEQ ID NO 41392; 3071pp + Sequence Listing; English.

AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC arctivity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC expression by rectifying mutations or deletions in a patient's genome CC expression by rectifying mutations or deletions in a patient's genome CC expression by rectifying mutations or deletions in a patient's genome CC expression by the patients own production of (I). Additionally, (I) CC the nucleic acids into a host cell and culturing the cell to express the CC polynucleotides may be used to provent and culturing the cell to express the CC diagnose and treat immune/haematopoietic-related diseases, especially to AAK67694 represent human immune/haematopoietic antigen genomic CC sequences from the present invention. AAK64703 to AAK64703 and AAM82169 crepresent sequences used in the exemplification of the present invention.

Sequence 99 BP; 18 A; 30 C; 29 G; 22 T; 0 other;

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RESULT 4
AAK86572
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                                                                                                                                                                      Matches
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Best Local :
07-NOV-2001 (first entry)
                                      AAK86572 standard; DNA; 99
                                                                                            4159 TCAGCCTCCCAAAGTGCTGGGATTACAAGCGTGAGCCA 4196
                                                                                                                    AAK86572;
                                                                                  62 TCAGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCA 99
                                                                                                                                                                   87;
                                                                                                                                                                           1.2%;
Similarity 88.8%;
                                                                                                                                                                   Conservative
                                                                                                                                                                0; Mismatches
                                                                                                                                                                      Score 80.4; DB 22; Pred. No. 1.5e-06;
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001)	piens. 7182-A2	immune/ha immune; iatic; ge
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2001WO-US01354. 2000US-0119065. 2000US-018664. 2000US-0186350. 2000US-0186350. 2000US-0186350. 2000US-0186350. 2000US-0198123. 2000US-0215186. 2000US-0216880. 2000US-0216880. 2000US-0216880. 2000US-0216880. 2000US-0216880. 2000US-0224518. 2000US-0224518. 2000US-0224518. 2000US-022575. 2000US-022566. 2000US-022566. 2000US-022575. 200U		/haematopoietic e; haematopoiet gene therapy; v
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		n genomic une/haema metastas
		i t
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299	27 29 29 29	14 21 21 25 25 25 26
29-SEP-2000; 200 29-SEP-2000; 200 02-OCT-2000; 200 03-NOV-2000; 200 08-NOV-2000; 200 08-NOV	-SEP- -SEP- -SEP-	SEP-SEP-SEP-SEP-SEP-SEP-SEP-SEP-SEP-SEP-
2000): 2000):	2000; 2000; 2000; 2000;	2000; 2000; 2000; 2000; 2000; 2000;
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18-0236370 18-0236380 18-0236802 18-0237037 18-0237039 18-0237039 18-0237039 18-0241785 18-0241785 18-0241785 18-0241785 18-0241808 18-0241808 18-0241808 18-0246476 18-0246471 18-0246475 18-0246475 18-0246475 18-0246475 18-0246475 18-0246525 18-024	236327	33065 34223 34274 34997 34998 35484 35834
888412111111111111111111111111111111111		

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amino acid sequences given in AAM92170 to AAM91921. [1] have cytostatic contivity, and can be used in gene therapy and vaccine production. (I) components and polynucleotides may be used in the prevention, diagnosis and creatment of diseases associated with inappropriate (I) expression. For cerample, they may be used to treat disorders associated with decreased cerample, they may be used to treat disorders associated with decreased that affect the activity of (I) by expressions in a patient's genome contact the activity of (I) by expressing inactive proteins or to polynucleotides may be used to product the finally, (I) concleic acids into a host cell and culturing the cell to express the concleic acids into a host cell and culturing the cell to express the concerts and cancer metastases of haematopoietic-derived cells. AAK64703 concerts and cancer metastases of haematopoietic antigen genomic concerts and cancer metastases of hae
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AAS32628/c
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Best Local :
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                                                                                                               Human; endocrine antigen; ds; cytostatic; antiinfertility; antidiabetic; thyroid-active; adrenal-active; androgenic; gastric; gene therapy; antibody; endocrine disorder; hormone imbalance; reproductive disorder; endocrine cancer; pancreatic disorder; endocrine cancer; pancreatic disorder; adrenal gland disorder; hirsutism; thyroid disorder;
                                                                                                                                                                                                                                                                                                                                                                                 4170 AAGTGCTGGGATTACAAGCGTGAGCCACTGCACC 4203
                                                                                                                                                                                                                                                                                                                                                                                                                                        08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
                                                                                                                                                                                                    Human genomic DNA for novel endocrine antigen, SEQ ID No 582.
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                                                                            Homo sapiens.
                                                                                                        hyperthyroidism;
                                                                                                                                                                                                                                    17-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                             AAS32628 standard; DNA; 98 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 99 BP;
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                                                                                                                                                                                                                                                                                                                                                                  AAGTGCTGGGATTACAGGTGTGAGCCACAGTGCC 99
                                                                                                                                                                                                                                                                                                                                                                                                                         951 to AAK64702 encode the human immune/haematopoietic antigen (I) acid sequences given in AAM82170 to AAM91921. (I) have cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0251989.
2000US-0251990.
2000US-0254097.
2000US-0259678.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                     hypothalamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 33 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruben SM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Pred. No. 2.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3071pp + Sequence Listing; English.
                                                                                                  disorder; vanishing testes syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
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18-AUG-2000
22-AUG-2000
22-AUG-2000
22-AUG-2000
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30-AUG-2000
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01-SEP-2000
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06-SEP-2000
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08-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
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27-SEP-2000;
29-SEP-2000;
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21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
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24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
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12-SEP-2000;
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28-JUN-2000;
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19-MAY-2000;
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                      2000US-0235484.
2000US-0235834.
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2000US-0236327.
2000US-0236367.
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2000US-0232397.
2000US-0232398.
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2000US-0229287.
2000US-0229343.
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2000US-0225759.
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2000US-0227182.
2000US-0227009.
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2000US-0234998
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2000US-0234274
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2000US-0231244
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2000US-0226681
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2000US-0225213.
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2000US-0220963
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2000US-0217496
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2000US-0205515
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08 NOV 2000;
01 NOV 2000;
17 NOV 2000;
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13-0CT-2000;
20-0CT-2000;
                                                                                                         17-NOV-2000
11-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
Isolated polypeptide for treating, preventing and prognosing disorders
                       WPI; 2001-457726/49.
                                                                      (HUMA-) HUMAN GENOME SCI INC.
                                              Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000US-0236802.
2000US-0236802.
2000US-0237037.
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2000US-0246525.

2000US-0246526.

2000US-0246527.

2000US-0246528.

2000US-0246528.

2000US-0246639.

2000US-0246639.

2000US-0246610.
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2000US-024617.

2000US-0246474.

2000US-0246475.

2000US-0246477.

2000US-0246477.

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2000US-0246478.
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2000US-0246613.
2000US-0249207.
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2000US-0249213.
2000US-0249214.
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2000US-0251869.
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                                               Ruben SM;
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RESULT 6
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CC antigen of the invention.
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   Matches
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Best Local 9
                              04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         related to the endocrine system including endocrine disorders, reproductive disorders, and gastrointestinal disorders and also for testing and detection e.g. diagnosis – \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID No 582; 558pp; English.
                                                                                                                                                                                                                                                                         Human; digestive system antigen; gene therapy; cancer; appendicitis; ulcerative colitis; infection; Hirschsprung's disease; chronic colitis; digestive system disorder; Meckel's diverticulum; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 98 BP; 22 A; 28 C; 29 G; 19 T; 0 other;
   18-APR-2000;
19-MAY-2000;
                                                                                                                                                                                                               WO200155314-A2
                                                                                                                                                                                                                                                                                                                                                                          05-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAK89189 standard; DNA; 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                 17-JAN-2001;
                                                                                                                                                                                 02-AUG-2001
                                                                                                                                                                                                                                                                                                                                       Human digestive system antigen genomic sequence
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2000US-0180628.
2000US-0184664.
2000US-0186350.
2000US-0189874.
2000US-019076.
2000US-0198123.
2000US-0205515.
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                                                                                                                                                 2001WO-US01324.
                                                                                                                 2000US-0179065
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Pred. No. 3e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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20-OCT 2000
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01-NOV 2000
08-NOV 2000
017-NOV 2000
17-NOV 2000
17-NO
The present invention provides the protein and coding sequences of number of human digestive system antigens. These can be used in the diagnosis, treatment and prevention of digestive system disorders, including cancer, Meckel's diverticulum, bacterial or parasitic infections, appendicitis, Hirschsprung's disease, chronic colitis
                                                                                                                                                                                                                                                                                             Polynucleotides encoding digestive system antigens, useful for diagnosing, treating, preventing and/or prognozing disorders of the digestive system, particularly cancer and cancer metastases -  \frac{1}{2} \int_{-\infty}^{\infty} \frac{1}{2}
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                                                                                                                                                                                                                                     Disclosure;
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                                                                                                                                                                                                                          SEQ ID NO 2765; 986pp; English.
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2000US-0241786.
2000US-0241808.
2000US-0241809.
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2000US-0254939.
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2000US-0251889.
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2000US-0254939.
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07-JUN-2000
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11-JUL-2000
11-JUL-2000
11-JUL-2000
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11-AUG-2000
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2000US-022514 2000US-022544 2000US-0215880 2000US-0216647 2000US-0217496 2000US-0217496 2000US-0217496 2000US-0220963 2000US-0220963 2000US-0225511 2000US-0225211 2000US-0225211 2000US-0225266 2000US-0225266 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-022527 2000US-0225447 2000US-022575 2000US-022575 2000US-022575 2000US-022575 2000US-02258681 2000US-02258681 2000US-0225928 2000US-0225928 2000US-0225928 2000US-0225928 2000US-02259344 2000US-0229345 2000US-0229344 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343 2000US-0229343

2000US -0231242 2000US -0231243 2000US -0231413 2000US -0231414 2000US -0231414 2000US -0232080 2000US -0232080 2000US -0232081 2000US -0232081 2000US -0232081 2000US -0232398 2000US -0232398 2000US -0232398 2000US -0232401 2000US -023401 2000US -023401

2000US-0233064 2000US-023405 2000US-0234274 2000US-0234997 2000US-023498 2000US-0235834 2000US-0235834 2000US-0235836 2000US-0236367 2000US-0236367 2000US-0236369 2000US-0236370 2000US-0236370 2000US-0236370 2000US-0237039 2000US-0237039

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RESULT 7
AAK89190
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Matches 86
18-APR-2000
19-MAY-2000
07-JUN-2000
30-JUN-2000
07-JUL-2000
07-JUL-2000
11-JUL-2000
14-JUL-2000
14-JUL-2000
14-AUG-2000
22-AUG-2000
22-AUG-2000
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24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ulcerative colitis. The present sequence is a genomic DNA fragment encoding a digestive system antigen of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; digestive system antigen; gene therapy; cancer; appendicitis; ulcerative colitis; infection; Hirschsprung's disease; chronic colitis; digestive system disorder; Meckel's diverticulum; ds.
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ilarity 87.8%;
Conservative (
               2000US-0179065.
2000US-0186628.
2000US-01866350.
2000US-0198123.
2000US-0198123.
2000US-0209467.
2000US-0214886.
2000US-0215135.
2000US-0216887.
2000US-0216887.
2000US-0217487.
2000US-0217487.
2000US-0217487.
2000US-0217487.
2000US-0217487.
2000US-0217487.
2000US-0217487.
2000US-02217487.
2000US-02252683.
2000US-0225256.
2000US-0225277.
2000US-0225277.
2000US-02257758.
2000US-02257758.
2000US-02257758.
2000US-02257758.
2000US-02257758.
2000US-02257758.
2000US-02257758.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 78.8; DB 22; Pred. No. 3e-06; 0; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
      22-AUG-2000

23-AUG-2000

30-AUG-2000

01-SEP-2000

01-SEP-2000

01-SEP-2000

01-SEP-2000

05-SEP-2000

06-SEP-2000

06-SEP-2000

08-SEP-2000

08-SEP-2000
   14-SEP-2000
14-SEP-2000
14-SEP-2000
14-SEP-2000
14-SEP-2000
21-SEP-2000
25-SEP-2000
25-SEP-2000
27-SEP-2000
29-SEP-2000
29-SEP-2000
29-SEP-2000
29-SEP-2000
29-SEP-2000
20-OCT-2000
02-OCT-2000
03-OCT-2000
03-OCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-SEP-2000;
14-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0227182.
2000US-0227009.
2000US-0228924.
2000US-0229287.
2000US-0229343.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000US-0234274.
2000US-0234997.
2000US-0234998.
2000US-0235484.
2000US-0235836.
2000US-0235836.
2000US-0235836.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-0232398.
2000US-0232399.
2000US-0232400.
2000US-0232401.
2000US-0233063.
2000US-0233065.
2000US-0233065.
2000US-0233065.
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2000US-0230437.
2000US-0230438.
                                                                                                                                                                                                                                                                                                                             2000US-0237038.
2000US-0237039.
2000US-0237040.
2000US-0239935.
2000US-0239937.
2000US-024960.
2000US-0241781.
2000US-0241785.
2000US-0241787.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0236367.
2000US-0236368.
2000US-0236369.
2000US-0236370.
2000US-0236802.
2000US-0237037.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0231413.
2000US-0231414.
2000US-0232080.
2000US-0232081.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0229344.
2000US-0229345.
2000US-0229509.
   2000US-0246528.

2000US-0246532.

2000US-0246609.

2000US-0246610.

2000US-0246611.

2000US-0246613.

2000US-0249208.
                                                                                                                       2000US-0246525.
2000US-0246526.
2000US-0246527.
                                                                                                                                                                2000US-0246474.
2000US-0246475.
2000US-0246476.
2000US-0246477.
2000US-0246477.
2000US-0246478.
2000US-0246523.
2000US-0246524.
                                                                                                                                                                                                                                                                   2000US-0241808.
2000US-0241809.
2000US-0241826.
2000US-0244617.
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2000US-0232397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0231242
2000US-0231243
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AAK85116/c
ID AAK85116 standard; L
XX AAK85116;
AC AAK85116;
XX
DT 07-NOV-2001 (first
                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-NOV-2000;
10-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences of a number of human digestive system antigens. These can be used in the diagnosis, treatment and prevention of digestive system disorders, including cancer, Meckel's diverticulum, bacterial or parasitic infections, appendicitis, Hirschsprung's disease, chronic colitis or ulcerative colitis. The present sequence is a genomic DNA fragment encoding a digestive system antigen of the invention.
                                                                                                                                                                                         4150
                                                                                                                                                                                                                                                           4090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polynucleotides encoding digestive system antigens, useful for diagnosing, treating, preventing and/or prognozing disorders of the digestive system, particularly cancer and cancer metastases - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                             Sequence 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 2766; 986pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME
                                                                                                                                                         61
                                                                                                                                                                       CCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACAAG
                                                                                                                                                                                                                                     GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
                                                                                                                                                      CCACCCGCCTTGGCCTCCCAAAGTGCTGGCATTACAGG
                                                                                                                                                                                                                       GTAGAGACGGGGTTTCACCATGTTGGCCAGGCTGGTCTCGAACTCCTGACCTCAGGTGAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-502630/55.
                                                                                                                                                                                                                                                                                           6;
                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barash SC,
                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                          BP; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0250391.

2000US-0251988.

2000US-0251988.

2000US-0256719.

2000US-0251479.

2000US-0251869.

2000US-0251869.

2000US-0251869.

2000US-0251869.

2000US-0251989.

2000US-0251989.

2000US-0251989.

2000US-0251989.

2000US-0251989.

2000US-0251989.
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2000US-0249264.
2000US-0249265.
2000US-0249297.
2000US-0249299.
2000US-0249299.
2000US-0249300.
2000US-0250160.
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2000US-0249217.
2000US-0249218.
2000US-0249244.
     (first entry)
                                                                                                                                                                                                                                                                                                                                                          A; 30 C; 28 G;
                                                                                                                                                                                                                                                                                                         1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SCI INC
                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                         Score 78.8; DB 22; Pred. No. 3e-06;
                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                      22 T; 0 other;
                                                                                                                                                                                     4187
                                                                                                                                                                                                                                                                                       12;
                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                       0;
26-JUL-2000

14-AUG-2000

12-AUG-2000

12-AUG-2000

22-AUG-2000

22-AUG-2000

22-AUG-2000

22-AUG-2000

23-AUG-2000

23-AUG-2000

25-SEP-2000

01-SEP-2000

01-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31 JAN 2000

04 FEB 2000

24 FEB 2000

02 MAR 2000

16 MAR 2000

17 MAR 2000

17 MAR 2000

19 MAY 2000

07 JUN 2000

28 JUN 2000

28 JUN 2000

07 JUL 2000

07 JUL 2000

07 JUL 2000

01 JUL 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immune/haematopoietic antigen genomic sequence SEQ ID NO:39928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune; haematopoietic;
             2000US-0232081.
2000US-0231968.
2000US-0232397.
2000US-0232398.
2000US-0232398.
2000US-0232400.
2000US-0232401.
2000US-0233403.
                                                                                                                                             2000US-0231244.
2000US-0231244.
2000US-0231413.
2000US-0231414.
2000US-0231414.
2000US-0232080.
                                                                                                                                                                                                                                                                                                  2000US-0226681.
2000US-02276868.
2000US-0227182.
2000US-0227009.
2000US-022924.
2000US-0229287.
2000US-0229343.
2000US-0229344.
2000US-0229344.
2000US-0229345.
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2000US-0215647.

2000US-0216647.

2000US-0216880.

2000US-0217487.

2000US-02188290.

2000US-02188290.

2000US-022964.

2000US-0224518.

2000US-0224518.

2000US-0225213.

2000US-0225213.

2000US-0225216.

2000US-0225216.
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                                                                                                                                                                                                                                                   2000US-0229513.
2000US-0230437.
2000US-0230438.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-0225268.
2000US-0225270.
2000US-0225447.
2000US-0225757.
2000US-0225757.
2000US-0225759.
2000US-0225759.
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2000US-0205515
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2000US-0190076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 etic; immune/haematopoietic antigen; cancer;
vaccine; metastasis; ds.
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2000US-0233065 2000US-0234274 2000US-0234997 2000US-0234997 2000US-0235484 2000US-02358834 2000US-0235834 2000US-0235836 2000US-0236367 2000US-0236367 2000US-0236369 2000US-0236369 2000US-0236369

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ACC XXX ACC XX
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ABA18411/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC amino acid sequences given in AAM82170 to AAM91921. (1) have cytostatic continuity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting CC the nucleic acids into a host cell and culturing the cell to express the CC protein. (I) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic derived cells. AAK64703 CC sequences from the present invention. AAK54942 to AAK5950 and AAA82169 CC sequences from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
05-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                     antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal;
antiparasitic; cardiant; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                    antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer;
antirheumatic; hepatotropic; cerebroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                  Human; nootropic; neuroprotective; cytostatic; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABA18411 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4173 TGCTGGGATTACAAGCGTGAGCCACTGCACCGGGC 4207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-483426/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC
16-AUG-2001.
                                                            WO200159063-A2
                                                                                                                        Homo sapiens.
                                                                                                                                                                                    neurological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nervous system related polynucleotide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAK54951
                                                                                                                                                                                                                                                                                                                                                immunosuppressive; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     35 TGCTGGGATTACAGGCATGAGCCACCGTGCCTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to AAK64702 encode the human immune/haematopoietic antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95 BP; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2%;
nilarity 88.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 39928; 3071pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-0251989.
2000US-0251990.
2000US-0254097.
2000US-0259678.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 26 C;
                                                                                                                                                                                       infection; nephrotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ruben SM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 77.4; DB 2:
Pred. No. 5.6e-06
D; Mismatches 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                    anti-HIV; antibacterial; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
                                                                                                                                                                                       gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 10742
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11-SEP-2000
21-SEP-2000
21-SEP-2000
25-SEP-2000
25-SEP-2000
26-SEP-2000
27-SEP-2000
29-SEP-2000
20-CCT-2000
20-CCT

2000US-0246524. 2000US-0246525. 2000US-0246526. 2000US-0246527.

2000US-0246609

2000US-0251030. 2000US-0251988. 2000US-0256719. 2000US-0251479. 2000US-0249212. 2000US-0249213. 2000US-0249214. 2000US-0249215. 2000US-0249215. 2000US-0249217. 2000US-0246611. 2000US-0246613. 2000US-0249207. 2000US-0249208. 2000US-0246474. 2000US-0246475. 2000US-0246476. 2000US-0246477.

2000US-0241808. 2000US-0241809. 2000US-0241826. 2000US-0244617. 2000US-0237040. 2000US-0239935. 2000US-0239937. 2000US-0240960. 2000US-0241221.

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The invention relates to novel genes (ABAl1004-ABA21534) and proteins (CC (ABB14678-ABB18001) useful for preventing, treating or ameliorating CC medical conditions e.g. by protein or gene therapy. The genes are CC isolated from a range of human tissues disclosed in the specification. CC The nucleic acids, proteins, antibodies and (anthagonists are useful CC in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast CC and ovarian cancer and other cancers of the adrenal gland, bone, bone CC marrow, breast, gastrointestinal tract, liver, lung, or urogenital; CC marrow breast, gastrointestinal tract, liver, lung, or urogenital; CC disease, multiple sclerosis, rhemmatoid arthritis and ulcerative CC disease, multiple sclerosis, rhemmatoid arthritis and ulcerative (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and CC epilepsy; and (f) infectious diseases such as viral, bacterial, fungal cancer infectious diseases such as viral, bacterial,
RESULT 10
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                         24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                antirheumatic; hepatotropic; cerebroprotective; antiinflammatory;
antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal;
antiparasitic; cardiant; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antisikling; antianaemic; antiarthritic; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human nervous system related polynucleotide SEQ ID NO 10746.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JAN-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                          neurological disease; infection; nephrotropic; gene therapy; vaccine;
                                                                                                                                                                                   17-JAN-2001;
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                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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The invention relates to novel genes (ABA11004-ABA21534) and proteins (ABB14678-ABB18001) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification
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17-NOV-2000;
17-NOV-2000;
                                                               Disclosure; SEQ
                                                                                                  Nucleic acids encoding useful for preventing,
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                                                                                       metastases
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                                                             ID NO 10746; 1701pp +
                                                                                                                                                                                   SCI INC.
                                                                                               3224 human
diagnosing
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                                                                                           nervous system antigen polypeptides, and/or treating nervous system
                                                          Sequence Listing; English
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RESULT 11
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Matches 85
30-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38
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                                                2000US-0184664
2000US-0186350
2000US-0189874
2000US-0190076
2000US-0198123
2000US-029515
2000US-0205515
2000US-0214886
2000US-0215135
2000US-0216647
2000US-0216647
2000US-0216880
2000US-0217488
2000US-0217487
2000US-0218290
2000US-02182963
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  2000US-0224518
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2000US-0180628
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); Mismatches
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Pred. No. 6.2e-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO: 661.
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14 - AUG - 2000; 14 - AUG - 2000;

2000US-0224519 2000US-0225213 2000US-0225264 2000US-0225266 2000US-0225268 2000US-0225269 2000US-0225270 2000US-022527447 2000US-0225757 2000US-0225759 2000US-0225759 2000US-0226681 2000US-0226681 2000US-0226681 2000US-0226681 2000US-0226681 2000US-0226681

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Query Match
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17-NOV-2000;
17-NO
                                                                                               The invention relates to an isolated polypeptide comprising an amino acid sequence at least 90% identical to 188 amino acid sequences fully defined in the specification and encoded by 188 cDNA clones fully defined in the specification. The invention also relates to a fragment having biological activity, a domain, an epicope, full length protein, variant, allelic variant or a species homologue of the fully defined sequence. The polynucleotide and polypeptide are useful for treating, preventing and/or prognosing disorders related to the pancreas including pancreatic cancer, pancreatitis, diabetes, endocrine disorders such as acromegaly or hyperthyroidism, and gastrointestinal disorders such as Crohn's disease and duodenal ulcers. The present sequence encodes a pancreatic cancer-related polypeptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated pancreatic cancer polypeptide for treating, preventing and/prognosing disorders related to the pancreas including pancreatic cancers and also for testing and detection e.g. diagnosis \dot{}
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 661; 537pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC
                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-457717/49.
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2000US-0246523.
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2000US-0251051.
2000US-0251050.
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                                                         A; 29 C;
     1.2%;
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                                                         30 G;
     Score 77.2;
                                                         21 T; 0 other;
          DB 22;
     Length 98
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or

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2000US-0230438
2000US-0230438
2000US-0231244
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08-NOV-2000;
08-NOV-2000;
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08-NOV-2000;

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2000US - 0244617.
2000US - 0246477.

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2000US-0227009. 2000US-0228924. 2000US-0229287. 2000US-0229343.

2000US-0229344 2000US-0229345

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RESULT 12
AAS32769
ID AAS32
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   11-JUL 2000

14-JUL 2000

26-JUL 2000

14-AUG 2000

14-AU
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04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
18-APR-2000;
19-MAY-2000;
07-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; endocrine antigen; ds; cytostatic; antiinfertility; antidiabetic; thyroid-active; adrenal-active; androgenic; gastric; gene therapy; antisense-therapy; antibody; endocrine disorder; hormone imbalance; reproductive disorder; endocrine cancer; pancreatic disorder; reproductive disorder; adrenal gland disorder; hirsutism; thyroid disorder; hyperthyroidism; hypothalamic disorder; vanishing testes syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human genomic DNA for novel endocrine antigen, SEQ ID No 723.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS32769 standard; DNA;
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2000US-0224518
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0; Mismatches 13;
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2000US-0229344.
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2000US-0232400
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17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
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05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                   The invention relates to cDNAs encoding novel human endocrine antigens or a fragment having biological activity, a domain, an epitope, full length protein, variant, allelic variant or a species homologue of the cDNA/antigen. The DNAs and polypeptides are useful for preventing, treating or ameliorating a medical condition when administered (e.g. by gene therapy or antisense-therapy). Identifying mutations in the genes coding for the antigens is useful for diagnosing a pathological condition or a susceptibility to a pathological condition. The DNAs, antigens and antibodies raised against the antigens useful for treating, preventing and/ or prognosing disorders related to the endocrine system or hormone imbalance or reproductive disorders, cancers of endocrine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                   tissues, disorders of the pancreas (e.g. diabetes mellitus), the adrenal glands (e.g. hirsutism), ovaries, the thyroid (e.g. hyperthyroidism), the hypothalamus and testes (e.g. vanishing testes syndrome), many examples of diseases and disorders are given in the specification. The present sequence is genomic DNA fragment form a gene encoding an endocrine antigen of the invention.

Note: The sequence data for this patent did not form part
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17-NOV-2000;
                 4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polypeptide for treating, preventing and prognosing disorders related to the endocrine system including endocrine disorders, reproductive disorders, and gastrointestinal disorders and also for testing and detection e.g. diagnosis
                                                                                                                                   Sequence 98
                                                                                                                                                                    ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                     of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID No 723; 558pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-457726/49
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                                                                                    Local
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 GTAGAGACGGGGTTTCACCATGTTGGGCAGGCTGGTCTCAAACTCCTGACCTCAGGTGAT 60
                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-0249213

2000US-0249213

2000US-0249215

2000US-0249215

2000US-0249216

2000US-0249218

2000US-0249218

2000US-0249244

2000US-0249264

2000US-0249264

2000US-0249267

2000US-0249297

2000US-0249297

2000US-0249297

2000US-0259180

2000US-02551866

2000US-0251866

2000US-0251866
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2000US-0251989.
2000US-0251990.
2000US-0254097.
2000US-0259678.
                                                                   Conservative
                                                                                                                                   BP; 18 A; 29 C; 30 G; 21 T; 0 other;
                                                                                   1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SM;
                                                                     Pred. No. 6.2
); Mismatches
                                                                                                  Score 77.2;
                                                                                 6.2e-06;
                                                                                                  DB 22;
                                                                                                  Length 98;
                                                                     Indels
                                                                     0
                                                                     Gaps
                                                                       0;
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RESULT 13
AAL04206/c
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      14-AUG-2000

18-AUG-2000

22-AUG-2000

22-AUG-2000

22-AUG-2000

23-AUG-2000

30-AUG-2000

01-SEP-2000

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01-SEP-2000

01-SEP-2000

05-SEP-2000

06-SEP-2000

08-SEP-2000

08-SEP-2000

08-SEP-2000
                                                                                                                                                                                                                                     26-JUL-2000;
26-JUL-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4150 CCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACAAG
                                                                                                                                                                                                           14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAL04206 standard; DNA; 98
                                                                                                                                                                                         14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                       16-MAR-2000;
17-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human reproductive system related antigen DNA SEQ ID NO:
                                                                                                                                                                                                                                                                                            14-JUL-2000;
                                                                                                                                                                                                                                                                                                       11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 CCGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reproductive system related antigen; reproductive system disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; ds.
2000US-0229345.
2000US-0229509.
2000US-0229513.
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2000US-0230438.
2000US-0231243.
2000US-0231243.
2000US-0231243.
2000US-0231244.
                                                                                                                                          2000US-0224518
2000US-0225213
2000US-0225213
2000US-0225214
2000US-0225266
2000US-0225268
2000US-0225268
2000US-0225270
2000US-0225447
2000US-0225757
2000US-0225757
2000US-0225758
2000US-0225758
2000US-0225758
2000US-0225758
2000US-0225758
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2000US-0217487.
2000US-0217496.
2000US-0218290.
2000US-0220963.
2000US-0220964.
                                                                                                     2000US-0226868.
2000US-0227182.
2000US-0227009.
2000US-0228924.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-US01339
                                                                                                                                                                                                                                                                                                                                                             2000US-0205515
2000US-0209467
                                                                           2000US-0229344
                                                                                   2000US-0229287.
2000US-0229343.
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                                                                                                                                                                                                                                                                                                                                                     2000US-0214886
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2000US-0189874.
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08-SEP-2000 08-SEP-2000 08-SEP-2000 12-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000

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RESULT 14
AAL05463
ID AAL05
XX
AC AAL05
XX
DT 21-NC
XX
HUMAR
XX
HUMAR
XX
HOMO
OS HOMO
OS HOMO
OS HOMO
XX
PD 02-AL
XX
PD 02-AL
PR 04-F1
PR 04-F1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 85; Conserv
  31-JAN-2000;
04-FEB-2000;
24-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-NOV-2000
17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
08-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4090 GTAGAGACAGGGTTTTGCCGTGTTGGCCGGGCTGGTCTCGAACTCTTGACCTCGGGTGAT 4149
                                                                                                                                                                                                                                                                                                           \label{thm:human: reproductive system related antigen; reproductive system disorder; \\ \text{cancer; gene therapy; } ds.
                                                                                                                                                                                                                                                                                                                                                                                       Human reproductive system related antigen DNA SEQ ID NO: 8151.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4150 CCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACAAG 4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be use in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding protein of the invention.
                                                                                                         17-JAN-2001; 2001WO-US01339
                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                          21-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL05463 standard; DNA;
                                                                                                                                                                                                               WO200155320-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 98 BP; 24 A; 27 C; 27 G; 20 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCGCCTGCCTTAGCCTCCCAAAGTGCTGGGATTACAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001-465570/50
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2000US-0179065.
2000US-0180628.
2000US-0184664.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 6894; 1297pp + Sequence Listing; English.
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2000US-0251856.
2000US-0251868.
2000US-0251869.
2000US-0251989.
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2000US-0249300.
2000US-0250160.
2000US-0250391.
2000US-0251030.
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2000US-0254097
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2000US-0256719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 77.2; DB 2
Pred. No. 6.2e-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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2000US-0231413
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2000US-0232081
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2000US-0249214

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20 - CCT - 2000
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20 - CCT - 2000
21 - CCT - 200

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02-MAR-2000

116-MAR-2000

117-MAR-2000

119-MAY-2000

07-JUN-2000

07-JUN-2000

07-JUL-2000

07-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-AUG-2000

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13-OCT-2000
20-OCT-2000
20-NOV-2000
20-NOV-2000
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20-NOV-2000
20-NOV-2000
20-NOV-2000
20-NOV-2000
21-NOV-2000
21-DEC-2000
20-DEC-2000
20-DEC
                                                                     Isolated is used i
   The
                                    Disclosure;
   present invention provides the protein and coding sequences
                                                                                                                        2001-465570/50.
                                                                                                                                                                                               HUMAN
                                                                     nucleic acid in preventing,
                                                                                                                                                         Barash SC,
                                    SEQ
                                                                                                                                                                                                                             2000US-0237040
2000US-023935
2000US-0239935
2000US-0241786
2000US-02411787
2000US-0241809
2000US-0244617
2000US-0246475
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2000US-0249211
2000US-0249218
2000US-0249288
2000US-0251869
2000US-0251869
2000US-0251869
2000US-0251989
2000US-0251989
                                                                                                                                                                                               GENOME
                                    ID NO
                                                                                                                                                                                               SCI
                                                                     molecule e, treating
                                8151; 1297pp + Sequence Listing; English
                                                                                                                                                             Ruben
                                                                                                                                                             MS
                                                                     encoding a reproductive system antigen
g or ameliorating a medical condition
     of a
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RESULT 15
AAK71848/c
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Best Local S
Matches 85
          30-JUN-2000
07-JUL-2000
11-JUL-2000
11-JUL-2000
11-JUL-2000
26-JUL-2000
26-JUL-2000
14-AUG-2000
                                                                                                                                                                                                                                                                                                                                                18-APR-2000;
19-MAY-2000;
07-JUN-2000;
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16-MAR-2000;
17-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immune/haematopoietic antigen genomic sequence SEQ ID NO:26660.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4148 ATCCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACA 4185
                                                                                                                                                                                                                                                                                                                                  07-JUN-2000;
28-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200157182-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; gene therapy; vaccine; metastasis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK71848 standard; DNA; 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-NOV-2001 (first entry)
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Similarity 86.7%;
85; Conservative
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2000US-0184664.
2000US-0186350.
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Pred. No. 6.2e-06;
0; Mismatches 13
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QΥ
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                             В
                                                                                                                                                                              CC AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (1) CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cartivity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally. (I) CC supplement the patients own production of (I). Additionally. (I) CC polynucleotides may be used to produce the secreted (I), by inserting CC the nucleic acids into a host cell and culturing the cell to express the CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK64703 cn present sequences used in the exemplification of the present invention.
                                                                                                       Query Match 1.2%; Score 77.2; DB 22
Best Local Similarity 86.7%; Pred. No. 6.2e-06;
                                                                                            Matches
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17-NOV-2000;
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17-NOV-2000;
4150 CCACCCACCTCAGCCTCCCAAAGTGCTGGGATTACAAG 4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
                                                                                                                                                   Sequence 98 BP; 24 A; 29 C; 26 G; 19 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 26660; 3071pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-483426/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA-) HUMAN GENOME SCI INC.
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                                                                                            0; Mismatches
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Search completed: January 4, 2003, 13:20:12 Job time: 834 secs